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Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 200250844-A 74 26-MAR-2002;
ISIS PRARMACEUTICALS INC
OS Unidentified
PD 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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Mixaglia, L.J., Worc, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 75 26-MAR-2002;
ISIS PHARMARENTICALS INC
OS Unidentified
N JP 2002508944-A/75
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JP 2002508944-A/75.
         0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 7.9e+02;
tive 0; Mismatches 0; Indels
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IG:232333093
JP 200250894-A/74.
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Antisense modulation of human MDM2 expression
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BD 2002508944-A/73
BD 20-MAR-2002
BD 26-MAR-1999 JP 2000538025
BP 26-MAR-1999 US 09/048810
BI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
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C12Q1/68,
C12N15/00
Strandedness: Single,
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Antieense modulation of human MDM2 expression FH
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Antisense modulation of human MDM2 expression FH
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Antisense modulation of human MDM2 expression Patent: JP 2002508944-A 72 26-MAR-2002; ISIS PHARMACEUTICALS INC OS Unidentified PN JP 2002508944-A/72 PD 26-MAR-2002 PF 26-MAR-1998 US 09/048810 PR 26-MAR-1998 US PR A6-MAR-1998 US PR 26-MAR-1998 US PR A6-MAR-1998 US 
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Antisense modulation of human MDM2 expression.
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Location/Qualifiers
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Location/Qualifiers

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PAT 18-SEP-2002

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unclassified.

1 (bases 1 to 20)
Miragila.L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 77 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/77
PD 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
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1 (bases 1 to 20)

Miraglia, L.J., Merco, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.

Autisense modulation of human MDM2 expression

Patent: JP 2002508944-A 78 26-MAR-2002;

ISIS PHARMACEUTICALS INC

S. Unidentified

PN JP 2002508944-A/78

PP 26-MAR-1200

PP 26-MAR-1999 US 09/048810

PR 26-MAR-1999 US 09/048810

PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX 1
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C12N1S/09,AG1K48/00,AG1P9/10,AG1P17/06,AG1P35/00,C07H21/04//
C1201/68,
C12N1S/00
Strandedness: Single;
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
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Location/Qualifiers
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100.0%; Pred. No. 7.9e+02;
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   TACCTACTGATGGTGCTGTA 1
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JP 2002508944-A/77.
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Best Local Similarity 100.C
Matches 20; Conservative
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BD138152/c
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E 1 (bases 1 to 20)

S Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression

L Patent: JP 2002508944-A 76 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/76
PD 26-MAR-2002
PF 26-MAR-199 UP 200538025
PR 26-MAR-199 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM, BRETT P MONIA,LEX M
26-WAR-2002
26-WAR-1999 JP 2000538025
26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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                                                                                              CI2N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
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C12Q1/68,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
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Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;
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"Antisense modulation of human MDM2 expression
L Patent: UP 2002508944-A 81 26-MAR-2002;
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PP 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
                          PAT 18-SEP-2002
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C12M15/00
Strandedness: Single;
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C12Q1/68,
C12N15/00
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
           20 bp DNA linear Antisense modulation of human MDM2 expression.
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JP 2002508944-A/80.
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TILLE Autisense modulation of human MNM2 expression
FITLE PARRAMCEUTICALS INC
FOURNAL ISP PARRAMCEUTICALS INC
FINT OS Unidentified
PN JP 2002508944-A/79
PD 20-ARAR-2002
PP 26-MAR-1999 JP --
PR 26-MAR-1999 JF --
PR 26-MAR
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26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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                    C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
C12N15/00
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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BD138153.1 GI:23233098
JP 2002508944-A/79.
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/organism='Unidentified'
Location/Qualifiers
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0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;

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                                                                                Strandedness: Single;
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AUTHORS TITLE JOURNAL REFERENCE

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B 1 (bases 1 to 20)
S Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
L Patent: JP 2002508944-A 83 26-MAR-2002;
SIS PHARMACEUTICALS INC
OS Unidentified
PN JP 200250844-A/83
PP 26-MAR-1999 JP 200538025
PP 26-MAR-1999 UP 200538025
PR 26-MAR-1999 UP 200538025
PR 26-MAR-1999 UP 200538029
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Miragila, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/84
PD 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 UP 2000538025
PR 26-MAR-1999 UP 2000538025
PR 26-MAR-1998 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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JP 2002508944-A/84
26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAWELA NERO, WARK J GRAHAM, BRETT P MONIA, LEX
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C12D1/68,
C12N15/00
Strandedness: Single,
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C12Q1/68,
C12N15/00
Strandedness: Single,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
13138158
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    .20
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ilarity 100.0%; Pred. No. 7.9e+02;
Conservative 0; Mismatches 0;
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/organism="unidentified"
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JP 2002508944-A/84.
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Best Local Similarity
Matches 20; Conserv
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BD138158/c
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1 (bases 1 to 20)

Miraglia L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
L. Patent: JP 2002508944-A 82 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/82
PP 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
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Topology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Topology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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BD138156.
BD138156.1 GI:23233101
JP 2002508944-A/82.
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Antiesense modulation of human MDM2 expression.
BD138157.1 GI:23233102
                                                                        /organism='Unidentified'
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/organism='Unidentified'
Location/Qualifiers
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0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;

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LOCUS
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BD138156/c
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Strandedness: Single;
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JP 2002508944-A/87.
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                                                                                                                                                                                                       JP 2002508944-A/85
26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/044810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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                                                              0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 7.9e+02;
ive 0; Mismatches 0; Indels
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                                                                                                                                                                                 linear
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Antisense modulation of human MDM2 expression.
BD138159
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Antisense modulation of human MDM2 expression.
BD138160
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Location/Qualifiers
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Strandedness: Single;
                                                                                                       450 ACTTATACTATGAAAGAGGT 469
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JP 2002508944-A/86.
                                                                                   Conservative
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1 (bases 1 to 20)
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unidentified
                                                           Query Match
Best Local Similarity
Matches 20; Conserv
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BD138160/c
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KEYWORDS
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AUTHORS
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JOURNAL
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KEYWORDS
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I (bases 1 to 20)

Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.

Antisense modulation of human MDM2 expression
Antisense modulation of human MDM2 expression

Antisense modulation of human MDM2 expression

ISIS PHARMACEUTICALS INC
OS Unidentified
DN JP 2002508944-A/87

PD 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 86 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/86
PP 36-MAR-1999 JP 200538025
PP 26-MAR-1999 US 09/048810
PP 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
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C12Q1/68,
C12N15/00
Strandedness: Single;
Topology: Linear;
Antisense modulation of human MDM2 expression FH Key
Location/Qualifiers
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C12Q1/68,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Location/Qualifiers
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JP 2002508944-A/88
26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1999 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
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1 (bases 1 to 20)

Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression

* Patent: JP 2002508944-A 89 26-MAR-2002;
ISIS PHARMACEUTICALS INC
                                                                                                                                                                                                                                     unclassified.

1 (bases 1 to 20)

Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 200208944-A 88 26-MAR-2002;
                                    Gaps
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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       Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 7.9e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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BD138163. B1:23233108
Up 2002508944-A/89. Unidentified unidentified unidentified unidentified
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Strandedness: Single;
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BD138162/c
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BD138163/c
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BD133164-1 GI:3233109

Up 2002508944-A/90.

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I (Dases 1 to 20)

Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.

Antiesnse modulation of human MDM2 expression

L Patent: JP 2002508944-A/90

SO Unidentified

NS Unidentified

NS Unidentified

SS Unidentified

PD 26-MAR-2002

PP 26-MAR-1999 JP 2000538025

PR 26-MAR-1999 US 09/048810

PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
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26-WAR-1999 JP 200538025
26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
                                                                26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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C12Q1/68,
C12N15/00
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C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
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Topology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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BD138164
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Location/Qualifiers
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/organism='Unidentified'
Location/Qualifiers
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7.9e+02;
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JP 2002508944-A/89
26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/04881
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Strandedness: Single;
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C12N15/00
Strandedness: Single;
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BD138167.1 GI:23233112
BD138167.1 GI:23233112
Unidentified
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BD138167/c
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TITLE
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E 1 (Dases 1 to 20)
S Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
L Patent: JP 2002508944-A 91 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/91
PD 26-MAR-2002
PF 26-MAR-1999 UP 200538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
                                                                                                                                                                                                                                                                                                                              26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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JP 2002508944-A/92
26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1999 US 09/048810
LOREN J MIRAGLIA, PAMBLA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
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1 (Dases 1 to 20)

Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 92 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/92
PD 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 UP 2000538025
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
BD138165
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Antisense modulation of human MDM2 expression.
BD138166
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/db_xref="taxon:32644"
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525 CAACATATTGTATATTGTTC 544
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JP 2002508944-A/91.
unidentified
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JP 2002508944-A/92.
                  20 CAACATATTGTATATTGTTC
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BD138165/c
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Partisense modulation of human MDM2 expression
Patent: JP 2002508944-A 93 26-MAR-2002;
SIST PHARMACEUTICALS INC
OS Unidentified
PD 26-MAR-2002
PD 26-MAR-1999 JP 2000538025
PF 26-MAR-1999 US 09/048810
PR 26-MAR-1998 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX P
C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
C12N15/08
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Antisense modulation of human MDM2 expression.
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Location/Qualifiers
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Location/Qualifiers
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20; Conservative 0; Mismatches
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    .20
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BD138171/c
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1 (bases 1 to 20)
Miraglia,L.J., Proc,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 95 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/95
PD 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1998 US 09/048810
PI 26-MAR-1998 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX N
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C12Q1/68,
C12N15/00
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CI2Q1/68,
CI2N15/00
                                                                                                                                           1 (bases 1 to 20)
Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Partisense modulation of human MDM2 expression
Pattisense modulation of human MDM2 expression
Pattisense modulation of human MDM2 expression
Pattisense modulation of human MDM2 expression
ISIS PHARMACEUTICALS INC
S Unique MAR-2002
PR 26-MAR-1999 JP 2000538025
PR 26-MAR-1998 US 09/048810
PI LOREN J MIRAGLIA, PAMMELA NERO, MARK J GRAHAM, BRETT P MONTA LEXT
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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                         20 bp DNA lin
Antisense modulation of human MDM2 expression.
BD138168
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                                                                       BD138168.1 GI:22233113
JP 2002508944-A/94.
unidentified
unidentified
unclassified.
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Best Local Similarity 100.
Matches 20; Conservative
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BD138169/c
RESULT 548
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JP 2002508944-A/96
26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1999 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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CI2N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
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            Topology: Linear; Antisense modulation of human MDM2 expression FH Location/Qualifiers
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Strandedness: Single;
Topology: Linear;
Artisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
BD138171
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/organism='Unidentified'
Location/Qualifiers
                                                              /organism='Unidentified'
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Strandedness: Single;
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Matches 20; Conservative
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unclassified.

1 (bases 1 to 20)
Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 99 26-MAR-2002;
SS Unidentified
BN JP 2002508944-A/99
BD 26-MAR-2002
PP 26-MAR-1999 UP 2000538025
PP 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
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Antisense modulation of human MDM2 expression FH
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BD138174
BD138174.1 GI:23233119
JP 2002508944-A/100.
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JP 2002508944-A/99.
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Location/Qualifiers
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BD138174/c
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BD138173/c
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JP 2005508944-A/98
26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
Strandedness: Single,
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
                                                             unclassified.

1 (bases 1 to 20)
Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 97 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/97
PD 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PR 10-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEY
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1 (bases 1 to 20)

Miragliab.L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 98 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PD 26-MAR-2002
PP 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 UP 2000538025
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
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/db_xref="taxon:32644"
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C12N15/00
Strandedness: Single;
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BD138171.1 GI:23233116
JP 2002508944-A/97.
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JP 2002508944-A/98.
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JP 2002508944-A/103.
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SSION BD138175

BD138
Miragila, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M. Autisense modulation of human MDM2 expression
Autisense modulation of human MDM2 expression
Patent: JP 2002508944-A 100 26-MAR-2002;
ISIS PHARMACEUTICALS INC
BY JOUGEOSE 2014
BY JOUGEOSE 2014
BY 26-MAR-1999 UP 2000538025
BR 26-MAR-1999 US 09/048810
BR 26-MAR-1999 US 09/048810
BR 26-MAR-1999 US 09/048810
BR 10-REN J MIRAGILA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
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C12Q1/68,
C12N15/00
Strandedness: Single;
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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PAT 18-SEP-2002
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                                                                               Gaps
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                                    ch 0.8%; Score 20; DB 1; Length 20; 1 Similarity 100.0%; Pred. No. 7.9e+02; 20; Conservative 0; Mismatches 0; Indels
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Antisense modulation of human MDM2 expression.
BD138177
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Antisense modulation of human MDM2 expression.
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Miraglia, L. J. Nero, P., Graham, M. J., Monia, B. P. and Cowsert, L. M. Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 105 26-MAR-2002;
ISTS PHRAMACEUTICALS INC
OS Unidentified
PD 26-MAR-2002
PD 26-MAR-2002
PF 26-MAR-1999 JP 2000538025
PF 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12N15/00
Strandedness: Single,
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Antisense modulation of human MDM2 expression.
BD138180 1 GI:23233125
JP 2002508944-A/106.
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JP 2002508944-A/105.
unidentified
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AUTHORS Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M. Patents modulation of human MDM2 expression
Fulls Antisense modulation of human MDM2 expression
Fulls Patent: JP 2002508944-A 104 26-MAR-2002;
SNT OS Unidentified
PN JP 2002508944-A/104
PD 26-MAR-1999 JP --
PR 26-MAR
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26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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                                                      26-MAR-1999 JP 2000538025
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26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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C12MIS/00
Strandedness: Single;
Topology: Linear;
Antisense modulation of human MDM2 expression FH Key
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
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C12N15/00
Strandedness: Single;
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JP 2002508944-A/103
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In (Dases 1 to 20)

Miraglial, L.J., Mero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.

Antisense modulation of human MDM2 expression

Disology

Disology

Unidentified

PN JP 2002508944-A/108

PN JP 2002508944-A/108

PN JP 2002508944-A/108

PN 26-MAR-1999 JP 2006538025

PR 26-MAR-1999 US 09/048810

PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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1 (Dases 1 to 20)
Miraglia,L.', Nero.P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 109 26-MAR-2002;
ISIS PHARMAGEUTICALS INC
S Unidentified
PR 2002508944-A/109
PD 26-MAR-1999 UP 200538025
PP 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 10AEN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
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C1201/68,
C12N15/00
Strandedness: Single,
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
source 1..20
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100.0%; Pred. No. 7.9e+02;
                                                  BD138182 20 bp DNA line
Antisense modulation of human MDM2 expression.
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Antisense modulation of human MDM2 expression.
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Conservative 0; Mismatches
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/organism="unidentified"
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JP 2002508944-A/109.
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BD138182/C
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LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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1 (bases 1 to 20)
Miragila.L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Fatent: JP 2002508944-A 107 26-MAR-2002;
Unidentified
PN JP 2002508944-A/107
PD 26-MAR-2002
PP 26-MAR-1999 JP 2005538025
PR 26-MAR-1998 US 09/048810
PR 26-MAR-1998 US 09/048810
PI LOREN J MIRAGILA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX P
                                             CI2NI5/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
C12N15/00
Strandedness: Single;
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Strandedness: Single;
Topology: Linear:
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;
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JP 2002508944-A/107.
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E 1 (Dases 1 to 20)
S Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
L Patent: JP 2002508944-A 110 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/110
PD 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM, BRETT P MONIA,LEX M
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26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, WARK J GRAHAM, BRETT P MONIA, LEX
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C12Q1/68,
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                        Topology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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0.8%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                              linear
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BD138184.
BD138184.1 GI:23233129
JP 2002508944-A/110.
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Strandedness: Single;
            Strandedness: Single;
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BD138185.1 GI:23233130
BD138185.1 GI:23233130
BD138185.1 GI:23233130
BD138185.1 GI:23233130
Unidentified
Unclassified.
SM unidentified
Unclassified.
B i (bases I to 20)
SM iragila, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
L Patent. JP 2002508944-A 111 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/111
PD 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 UP 2000538025
PR 26-MAR-1999 US 09/040810
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BD138186.1 GI:22233131
JP 2002508944-A/112.
unidentified
unclassified.
EMiragliat.L.1, Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
L Patent: JP 2002508944-A 112 26-MAR-2002;
SS Unidentified.
D 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 US 099/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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C12Q1/68,
C12N15/00
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression
Location/Qualifiers
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Strandedness: Single,
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BD138189/c
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AUTHORS
TITLE
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Antisense modulation of human MDM2 expression
Batent: JP 2002508944-A 113 26-MAR-2002;
S PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/113
PD 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR LOREN J MIRAGILA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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Strandedness: Single;
Topology: Linear;
Antisense modulation of human MDMZ expression FH
Location/Qualifiers
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                                                                                                       0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 7.9e+02; tive 0; Mismatches 0; Indels
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Location/Qualifiers
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BD138188/c
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unclassified.

1 (bases 1 to 20)
Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 114 26-MAR-2002;
ISIS PHARMAGEUTICALS INC
BN JP 2002508944-A/114
BD 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US DERO,MARK J GRAHAM,BRETT P MONIA,LEX |
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C12Q1/68,
C12N15/00
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C12Q1/68,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;
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BD138192/c
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BD138190.1 GI:23233135

PD 2002508944-A/116.

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I (bases 1 to 20)

Miraglia.Lu., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.

Antisense modulation of human MDM2 expression

L Patent: JP 2002508944-A 116 26-MAR-2002;

SOS Unidentified

PD 26-MAR-2002

PP 26-MAR-1999 JP 200538025

PR 26-MAR-1999 US 009/04810

PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
                                                                                                                                                                                                                                                                                                                                                                                  26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 (bases 1 to 20)
Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 117 26-WAR-2002;
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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                                     0.8%; Score 20; DB 1; Length 20; 00.0%; Pred. No. 7.9e+02; Ve 0; Mismatches 0; Indels
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Antieense modulation of human MDM2 expression.
BD138191
                                                                                                                                                                           20 bp DNA lin
Antieense modulation of human MDM2 expression.
BD138190
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                                   Query Match

0.8%; Score 20, 27.9

Best Local Similarity 100.0%; Pred. No. 7.9

Matches 20; Conservative 0; Mismatches
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Strandedness: Single;
                                                                                         787 CATCTAGAAGGAGGAATT 806
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                                                                                                        CATCTAGAAGGAGAGCAATT 1
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JP 2002508944-A/117.
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Matches 20; Conserv
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ACCESSION
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BD138190/c
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AUTHORS
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Miragila, L. J. Nero, P., Graham, M. J., Monia, B. P. and Cowsert, L. M. Antisense modulation of human MDM2 expression
Eatent: JP 2002508944-A 118 26-MAR-2002;
Sundentified
SN Unidentified
DN J 2002508944-A/118
PD 26-MAR-2002
PF 26-MAR-1999 UP 2000538025
PF 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGILA, PAMBLA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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                                                              26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
                                                                                                                                                                       C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//C12Q1/68,
C12D1/68,
C12N15/00
Strandedness: Single,
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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BD138192.
BD138192.1 GI:2323137
JP 20026/nPo.4.*/
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Location/Qualifiers
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Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;
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C12N15/00
Strandedness: Single,
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Unidentified
JP 2002508944-A/117
26-MAR-2002
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Matches 20; Conserv
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ACCESSION
VERSION
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BD138195/c
LOCUS
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SOURCE
ORGANISM
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TITLE
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Antisense modulation of human MDN2 expression
Patent: JP 2002508944-A 119 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/119
PP 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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SM unidentified
unclassified.
B (bases 1 to 20)
S Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
L Patent: JP 2002508944-A 120 26-MAR-2002;
OS Unidentified
PN JP 2002508944-A/120
PN JP 2002508944-A/120
PP 26-MAR-1999 JP 2000538025
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C12Q1/68,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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llarity 100.0%; Pred. No. 7.9e+02;
Conservative 0; Mismatches 0; Indels
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    Indels
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Antisense modulation of human MDM2 expression.
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  0; Mismatches
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                                        20 TTCAGATGAATTATCTGGTG 1
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JP 2002508944-A/120.
20; Conservative
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Best Local Similarity
Matches 20; Conserva
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VERSION
KEYWORDS
SOURCE
ORGANISM
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ACCESSION
VERSION
KEYWORDS
SOURCE
ORGANISM
                                                                                        RESULT 573
BD138193/c
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AUTHORS
TITLE
JOURNAL
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SM unidentified
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SM traglia, L.J., Norco, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
L. Patent: JP 2002508944-A 121 26-MAR-2002;
ISIS PHARACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/121
PD 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
                                                                  C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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llarity 100.0%; Pred. No. 7.9e+02;
Conservative 0; Mismatches 0; Indels
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Location/Qualifiers
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Antisense modulation of human MDM2 expression.
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Location/Qualifiers
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7.9e+02;
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Strandedness: Single;
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Strandedness: Single;
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BD138195.1 GI:23233140
JP 2002508944-A/121.
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DD138198

ON Antisense modulation of human MDM2 expression.

M BD138198

BD138198

BD138198

BD138198

I G1:23233143

I DP 2002508944-A/124.

SM unidentified

unclassified.

I (bases 1 to 20)

SM Antisense modulation of human MDM2 expression

L SIS PHARMACEUTICALS INC

OS Unidentified

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PR 26-MAR-1999 UP 200538025

PR 26-MAR-1999 US 2009538025

PR 26-MAR-1999 US 09/048810
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C12Q1/68,
C12N15/00
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                                              Topology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Best Local Similarity 100.0%; Pred. No. 7.9

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C12Q1/68,
C12N15/00
Strandedness: Single;
                                                                                                                                                                                                                                                                               880 CCCTTTCCTTTGATGAAGC 899
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BD138198/c
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BD138199/c
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TITLE
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unclassified.

In Clases 1 to 20)

S. Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antieense modulation of human MDM2 expression

L. Patent: JP 2002508944-A 123 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/123
PD 26-MAR-2002
PP 26-MAR-1999 UP 200538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
                                                                                                                                             unidentified
unclassified.

E 1 (bases 1 to 20)
S Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
L Patent: JP 2002508944-A 122 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/122
PD 26-MAR-1999
PP 26-MAR-1999 UP 200538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM, BRETT P MONIA,LEX M
                                                                                                                                                                                                                                                                                          26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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C12Q1/68,
C12N15/00
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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                                                   20 bp DNA line
Antisense modulation of human MDM2 expression.
BD138196
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Antisense modulation of human MDM2 expression.
BD138197
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JP 2002508944-A/123.
unidentified
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JP 2002508944-A/122.
unidentified
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DEFINITION
ACCESSION
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BD138197/c
LOCUS
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KEYWORDS
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                             RESULT 576
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AUTHORS
TITLE
JOURNAL
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Location/Qualifiers
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JP 2002508944-A/127.
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Best Local Similarity 100.0
Matches 20; Conservative
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Best Local Similarity 100.0
Matches 20; Conservative
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unclassified.
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DEFINITION
ACCESSION
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KEYWORDS
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BD138201/c
LOCUS
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BD138202/c
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TITLE
JOURNAL
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Patent: JP 2002508944-A 115 26-MAR-2002;
Sunidentified
PN JP 2002508944-A/125
PP 26-MAR-2002
PP 26-MAR-1999 UP 2000538025
PP 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 10-REN J MIRAGILA, PAMBLA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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unclassified.

E 1 (bases 1 to 20)
S Mixaglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
L Patent: JP 2002508944-A 126 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/126
PN JP 2002508944-A/126
PN JP 2005508949-A/126
PN 26-MAR-1999 UP 200538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM, BRETT P MONIA,LEX M
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26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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CI2Q1/68,
CI2NI5/00
Strandedness: Single;
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CI2Q1/68,
CI2N15/00
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
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            BD138199
BD138199.1 GI:23233144
JP 2002508944-A/125.
unidentified
unidentified
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BD138200/c
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DEFINITION
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Miragila, L. J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M. Antiagila, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M. Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 127 26-MAR-2002;
OS Unidentified
OS Unidentified
PN JP 2002508944-A/127
PD 26-MAR-1999 UP 2002508944-A/127
PP 26-MAR-1999 UP 2000538025
PR 26-MAR-1998 US 09/048810
PR 26-MAR-1998 US 09/048810
PR 10-REN J MIRAGILIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
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Strandedness: Single;
Stropology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
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Location/Qualifiers
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. 7.9e+02;
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100.0%; Pred. No. 7.9
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DEFINITION
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VERSION
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ACCESSION
VERSION
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ORGANISM
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BD138204/c
LOCUS
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SM unidentified
unclassified.

Losses Lto 20)
S Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
L Patent: JP 2005508944-A 128 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/128
PD 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
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BD138203

ACCESSION
Antitense modulation of human MDM2 expression.
ACCESSION
VERSION
BD138203.1 GI:23233148

KEYWORDS
JP 2002508944-A/129.
SOURCE
Unidentified
Unclassified.

REFERENCE
AUTHORS
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AUTHORS
ALTIE
ANTISENSE MODESO8944-A 129 26-MAR-2002;
SOURCE
JP 2002508944-A/129
PD 26-MAR-2002
PF 26-MAR-1999
PD 26-MAR-1007
PI 1777.2
PP 26-MAR-1007
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26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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C12Q1/68,
C12N15/00
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C12Q1/68,
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Antisense modulation of human MDM2 expression Location/Qualifiers
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Location/Qualifiers
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ilarity 100.0%; Pred. No. 7.9e+02
Conservative 0; Mismatches 0

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unidentified
unclassified.
1 (bases I to 20)
Miraglia,L.J., Mero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 130 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/130
PD 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 US 09/048810
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C12Q1/68,
C12N15/00
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1 (bases 1 to 20)
Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M. Antisense modulation of human MDM2 expression
                                                  Gaps
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Location/Qualifiers
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Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 7.9e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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JP 2002508944-A/131. unidentified unidentified
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JP 2002508944-A/130.
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1 (Dases 1 to 20)
Miraglia,L.', Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antiense modulation of human MDM2 expression
Patent: JP 2002508944-A 133 26-MAR-2002;
ISIS PHARMACEUTICALS INC
SU Unidentified
PN JP 2002508944-A/133
PD 26-MAR-1999 UP 200538025
PP 26-MAR-1999 UP 200538025
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 1-OREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
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C12Q1/68,
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Partisense modulation of human MDM2 expression
Patent: JP 2002508944-A 134 26-MAR-2002;
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PN JP 2002508944-A/134
PD 26-MAR-2002
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llarity 100.0%; Pred. No. 7.9e+02;
Conservative 0; Mismatches 0; Indels
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Strandedness: Single;
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JP 2002508944-A/133.
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26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
Strandedness: Single;
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Cl2XIS/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
Cl2Q1/68,
Cl2XIS/00
Strandedness: Single;
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Patent: JP 2002508944-A 132 26-MAR-2002;
OS Unidentified
PN JP 2002508944-A/132
PD 26-MAR-2002
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Location/Qualifiers
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Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;
Patent: JP 2002508944-A 131 26-MAR-2002;
ISIS PHARMACEUTICALS INC
S Unidentified
PN JP 2002508944-A/131
PD 26-MAR-2002

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JP 2002508944-A/132.
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ACCESSION VERSION KEYWORDS SOURCE ORGANISM

REFERENCE AUTHORS TITLE JOURNAL

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BD138206/c LOCUS RESULT 586

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DNA linear PAT 18-SEP-2002

NA Antisense modulation of human MDM2 expression.

NA BD138211

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BD18826-MAR-1999 US 09/048810

BD1 LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.

Antisense medulation of human MDM2 expression

Antisense medulation of human MDM2 expression

L Patent: JP 200350894-A 136 26-MAR-2002;

OS Unidentified

PN JP 2002508944-A/136

PN JP 2005508944-A/136

PN JP 2005508948-A/136

PN Z6-MAR-1999 US 005088025

PR Z6-MAR-1999 US 005080802

PR Z6-MAR-1999 US 005048810

PR LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM, BRETT P MONIA,LEX M
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
C12N15/00
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antieense modulation of human MDM2 expression.
BD138210 G1:23233155
JP 2002508944-A/136.
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TITLE
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TITLE
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26-MAR-1999 UP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
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1 (bases 1 to 20)
Miraglia,L.J., Nerco,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 135 26-MAR-2002;
Unidentified INC
OS Unidentified INC
PP 26-MAR-2002
PP 26-MAR-135
PP 26-MAR-1999 UP 200538025
PR 26-MAR-1999 UP 200538025
PR 26-MAR-1999 UP 200538029
PR 10RBN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX
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Antisense modulation of human MDM2 expression FH
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Antisense modulation of human MDM2 expression FH
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Antisense modulation of human MDM2 expression.
BD138209
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Location/Qualifiers
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Location/Qualifiers

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/mol_type="genomic DNA"
/db_xref="taxon:32644"

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    /organism="unidentified"

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/db_xref="taxon:32644"
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C12N15/00
Strandedness: Single,
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C12N15/00
Strandedness: Single,
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BD138209/c
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C12Q1/68,
C12N15/00
C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Topology: Linear,
Antisense modulation of human MDM2 expression FH
Location/Qualifier
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100.0%; Pred. No. 7.9e+02;
tive 0; Mismatches 0; Indels
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Location/Qualifiers
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/organism="unidentified"
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/db_xref="taxon:32644"

    .20
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    /mol_type="genomic DNA"
    /db_xref="taxon:32644"

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                C1201/68,
C12N15/00
Strandedness: Single;
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Best Local Similarity 100.0%
Matches 20; Conservative
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BD138212/c
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RESULT 593 BD138213/c

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Mirch 201

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C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
C12N15/00
Strandedness: Single;
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C12Q1/68,
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BD138213 20 bp DNA line
Antisense modulation of human MDM2 expression.
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/db_xref="taxon:32644"
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Strandedness: Single;
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JP 2002508944-A/139.
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Miragila, L., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M. Antisense modulation of human MDM2 expression
Antisense modulation of human MDM2 expression
Fatent: JP 2002508944-A 142 26-MAR-2002;
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C12NIS/00
Strandedness: Single;
Topology: Linear;
Antisense modulation of human MDM2 expression FH Key
Location/Qualifiers
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
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Antisense modulation of human MDM2 expression.
BD138217
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/organism='Unidentified'
Location/Qualifiers
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100.0%; Pred. No. 7.9e+02;
tive 0; Mismatches 0;

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/mol_type="genomic DNA"
/db_xref="taxon:32644"

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Strandedness: Single;
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Best Local Similarity 100.
Matches 20; Conservative
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unclassified.
E 1 (bases 1 to 20)
S Mizaglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
L Patent: JP 2002508944-A 141 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/141
PD 26-MAR-1999 JP 200538025
PR 26-MAR-1999 UP 200538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
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JP 2002508944-A/141
26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1999 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
C12N15/00
Strandedness: Single;
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Antisense modulation of human MDM2 expression FH Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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BD138216
BD138216.1 GT:23233161
JP 2002508944-A/142.
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Antisense modulation of human MDM2 expression.
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100.0%; Pred. No. 7.9e+02;
ative 0; Mismatches 0;

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JP 2002508944-A/141.
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BD138215/c
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BD138218.1 GI:33233163

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E 1 (bases 1 to 20)

Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.

Antisense modulation of human MDM2 expression

Antisense modulation of human MDM2 expression

ISIS PHARMACEUTICALS INC

OS Unidentified

PN JP 2002508944-A/144

PD 26-MAR-1999 JP 2000538025

PR 26-MAR-1999 JP 2000538025

PR 26-MAR-1999 US 09/048810

PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
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C12Q1/68,
C12N15/00
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1 (bases 1 to 20)
Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
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Topology: Linear;
Attisense modulation of human MDM2 expression FH
Location/Qualifiers
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llarity 100.0%; Pred. No. 7.9e+02;
Conservative 0; Mismatches 0; Indels
                                                             Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 7.9e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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BD138219
BD138219. GI:23233164
JP 2002508944-A/145.
unidentified
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BD138218
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Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 145 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/145
PD 26-MAR-1999 JP 200538025
PF 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Topology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
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Strandedness: Single;
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26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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                                                                              COWSERT
C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C1201/68,
C12N15/00
Strandedness: Single;
Topology: Linear;
Topology: Linear;
Antisense modulation of human MDM2 expression FH Key
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C12Q1/68,
C12N15/00
Strandedness: Single;
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Partis PRARMACEUTICALS INC
S Unidentified
PN JP 2002508944-A/149
PD 26-MAR-1998 JP 2000538025
PR 26-MAR-1998 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
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JP 2002508944-A/149.
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BD138223/c
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                                                                                                                                                                                                                                                               unidentified
unclassified.

In (bases 1 to 20)

Is Mixaglia,L.J., Mero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antieense medulation of human MDM2 expression

L Patent: JP 2002508944-A 147 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/147
PD 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
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LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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1 (bases 1 to 20)
Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M..
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 148 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/148
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
C12N15/00
Strandedness: Single;
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Antisense modulation of human MDM2 expression FH
        Length 20;
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100.0%; Pred. No. 7.98+02;
Live 0; Mismatches 0; Indels
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Antisense modulation of human MDM2 expression.
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Antisense modulation of human MDM2 expression.
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   0.8%; Score 20; DB 1; Le
100.0%; Pred. No. 7.9e+02;
ative 0; Mismatches 0;
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/organism="unidentified"
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/db_xref="taxon:32644"
                                                                 1105 GACAAGAACTCTCAGATGAA 1124
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JP 2002508944-A/147.
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JP 2002508944-A/148.
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BD138222/c
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C12Q1/68,
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/organism="unidentified"
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/organism="unidentified"
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Strandedness: Single;
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BD138226/c
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Antisense modulation of human MDM2 expression
Batent: JP 2002508944-A 150 26-MAR-2002;
Sunidentified
So Unidentified
BN J 2002508944-A/150
PP 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR LOREN J MIRAGILA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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D 2002508944-A 151 26-MAR-2002;
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D 2002508944-A 151 26-MAR-2002;
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26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
                                                                                                                                                          linear
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Antisense modulation of human MDM2 expression.
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    .20
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    /mol_type="genomic DNA"
    /db_xref="taxon:32644"

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20 TATATCAAGTTACTGTGTAT 1
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JP 2002508944-A/150.
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Best Local Similarity 100.0%
Matches 20; Conservative
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ACCESSION
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SOURCE
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BD138224/c
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E 1 (bases 1 to 20)

S Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.

Antisense modulation of human MDM2 expression

Antisense modulation of human MDM2 expression

I SIS PHARMACEUTICALS INC

OS Unidentified

PN JP 2002508944-A152 26-MAR-2002;

OS Unidentified

PN JP 2002508944-A152

PP 26-MAR-1999 JP 2000538025

PR 26-MAR-1999 UP 2000538025

PR 26-MAR-1999 US 09/048810

PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M ö ö PAT 18-SEP-2002 CI2NIS/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//C1201/68,
C12NIS/00
Strandedness: Single;
Topology: Linear;
Antisense modulation of human MDM2 expression FH Key Location/Qualifician Gaps ö .. 0 Topology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 7.9e+02; Matches 20; Conservative 0; Mismatches 0; Indels 0.8%; Score 20; DB 1; Length 20; .00.0%; Pred. No. 7.9e+02; linear Antisense modulation of human MDM2 expression. BD138226. BS28226.1 GI:23233171 ce 1. 20 /organism='Unidentified' Location/Qualifiers /organism='Unidentified'

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E 1 (bases 1 to 20)

S Miraglia, L. D., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antrisense modulation of human MDM2 expression

Antrisense modulation of human MDM2 expression

L Patent: JP 2002508944-A 153 26-MAR-2002;

ISIS PHARM-CEVITCALS INC

OS Unidentified

PN JP 2002508944-A/153

PD 26-MAR-1999 JP 200538025

PR 26-MAR-1999 US 09/048810

PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
                                                                                                                                                                                                                                                                                                     26-WAR-1999 JP 2000538025
26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
                    PAT 18-SEP-2002
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C12N1S/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
C12N1S/00
Strandedness: Single;
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C12Q1/68,
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1 (bases 1 to 20)
Miragila,L.J., Nerc,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 154 26-MAR-2002;
SIS PHARMAGEUTICALS INC
OS Unidentified INC
PD 26-MAR-2002
PD 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 009/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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           20 bp DNA line
Antisense modulation of human MDM2 expression.
BD138227
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Antisense modulation of human MDM2 expression.
BD138228
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Location/Qualifiers
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0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;
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                                                                     BD138227.1 GI:23233172 JP 2002508944-A/153. unidentified
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JP 2002508944-A/154.
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BD138227/c
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C12Q1/68,
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Topology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antieense modulation of human MDM2 expression.
BD138229
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100.0%; Pred. No. 7.9e+02;
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Antisense modulation of human MDM2 expression.
BD138230. GI:23233175
                                                                                                              'organism='Unidentified'
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/mol_type="genomic DNA"
/db_xref="taxon:32644"

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    /organism="unidentified"

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BD138230/c
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Matches 20; Conservative
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                               SM unidentified
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L (bases 1 to 20)

Miraglia,LJ., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression

L Patent: JP 2002508944-A 156 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/156
PD 26-MAR-2002
PP 26-MAR-199 UP 200538025
PR 26-MAR-199 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM, BRETT P MONIA,LEX M
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JP 2002508944-A/157.
Unidentified
I (bases 1 to 20)
S Miraglia,L.J., Mero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
I Patent: JP 2002508944-A 157 26-MAR-2002,
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/157
PD 26-MAR-2002
PP 26-MAR-1998 UP 200538025
PR 26-MAR-1998 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
                                                                                                                                                                                             26-WAR-1999 JP 2000538025
26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
C12N15/00
Strandedness: Single,
Topology: Linear;
Antisense modulation of human MDM2 expression FH Key
Location/Qualifiers
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C12Q1/68,
C12N15/00
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
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    /organism='Unidentified'

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Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;
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Miraglia, L. J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M. Antiaglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M. Antiagense modulation of human MDM2 expression
Patent: JP 2002508944-A 158 26-MAR-2002;
S. Unidentified
BN JP 2002508944-A/158
PD 26-MAR-2002
PP 26-MAR-1999 JP 200538025
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1998 US 09/048810
PR 10REN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
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                                                                                               Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 7.9e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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Strandedness: Single;
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1. .20
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                                                20; Conservative
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C12N15/00
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Fates and modulation of human MDM2 expression
Fatent: JP 2002508944-A 160 26-MAR-2002;
SS Unidentified
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SS Unidentified
FOR UNIDENTIALS INC
SC WAR-1999 JP 2000538025
FOR 26-MAR-1999 UP 2000538025
FOR 26-MAR-1999 US 09/048810
FOR 26-MAR-1999 US 09/048810
FOR 26-MAR-1999 US 09/048810
FOR 26-MAR-1999 US 09/048810
Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
Batent: JP 2002508944-A 159 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/159
PD 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX 1
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C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
C12N15/00
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C12Q1/68,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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JP 2002508944-A/160.
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/organism='Unidentified'
Location/Qualifiers

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Strandedness: Single;
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1 (bases 1 to 20)
Miraglia,L.J., Nerco,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human WDM2 expression
Patent: JP 2002508944-A 161 26-MAR-2002;
ISIS PHARMAGEUTICALS INC
S Unidentified
PN JP 2002508944-A/161
PD 36-MAR-1999 JP 200538025
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
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NALtsense modulation of human MDM2 expression.

NALtsense modulation of human MDM2 expression.

BD138236.1 G1:23233181

SD02508944-A/162.

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Unclassified.

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Antisense modulation of human MDM2 expression

L Patent: JP 2002508944-A 162 26-MAR-2002;

ISIS PHARMACEUTICALS INC

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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 7.9e+02; ive 0; Mismatches 0; Indels
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L (Dases 1 to 20)

Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense medulation of human MDM2 expression
Antisense medulation of human MDM2 expression

L Patent: JP 2002508944-A 165 26-MAR-2002;

ISIS PHARMACEUTICALS INC
ON Unidentified
PN JP 2002508944-A/165
PD 26-MAR-1099 JP 200538025
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
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DP 2002508944-A/165
26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1999 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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1 (bases 1 to 20)
Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 164 26-MAR-2002;
SIS PHARMAGEUTICALS INC
DN JC 2002508944-A/164
PD 26-MAR-2002
PP 26-MAR-1999 UP 200538025
PP 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 10-REN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX 1
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C12Q1/68,
C12N15/00
Strandedness: Single,
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Antisense modulation of human MDM2 expression
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Antisense modulation of human MDM2 expression.
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           26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12N15/00
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Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 163 26-MAR-2002;
S Unidentified
PN JP 2002508944-A/163
PD 26-MAR-2002
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Antisense modulation of human MDM2 expression FH
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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C12N15/00
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2002508944-A/162
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JP 2002508944-A/163.
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Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.

Antisense modulation of human MDM2 expression

L Patent: JP 2002508944-A 167 26-MAR-2002;

ISIS PHARWACEUTICALS INC

OS Unidentified

PN JP 2002508944-A/167

PD 26-MAR-1999 JP 200538025

PF 26-MAR-1999 US 09/048810

PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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E 1 (bases 1 to 20)
S Mirsglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
L Patent: JP 2002508944-A 168 26-MAR-2002;
ISIS PHARMACEUTICALS INC
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PN JP 2002508944-A/168
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PP 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
                                                    PAT 18-SEP-2002
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C12Q1/68,
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26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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Antisense modulation of human MDM2 expression FH
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Antisense modulation of human MDM2 expression.
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Location/Qualifiers
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/organism="unidentified"
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E 1 (bases 1 to 20)
S Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
L Patent: JP 2002508944-A 166 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/166
PD 26-MAR-2002
PP 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM, BRETT P MONIA,LEX M
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26-MAR-2002
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LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
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/organism='Unidentified'
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/organism="unidentified"
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Location/Qualifiers
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                                                                                                                                           26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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                                                                                                                                                                                                                     CI2NIS/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
CI2Q1/68,
CI2NI5/00
unclassified.

1 (bases 1 to 20)

Miraglia,L.J., Werc,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 170 26-MAR-2002;
SS PHARMACEUTICALS INC
OS JU 2002508944-A/170
PD 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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1 (bases 1 to 20)
Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Partisense medulation of human NDM2 expression
Patent: JP 2002508944-A.173 26-MAR-2002;
S Unidentified INC
ON Unidentified PN
P 2002508944-A/173
PD 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 US 09/048810
PP 10-REN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX N
                                                                                                                                                                              CI2N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
C12N15/00
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C12Q1/68,
C12N15/00
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/organism='Unidentified'
Location/Qualifiers
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Location/Qualifiers
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JP 2002508944-A/172
26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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                                                                                                                                                                                                                                                                                                                     unidentified
unclassified.
1 (bases 1 to 20)
Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 172 26-MAR-2002;
ISIS PHARMACEUTICALS INC
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Pred. No. 7.9e+02;
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Antisense modulation of human MDM2 expression.
BD138246
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Antieense modulation of human MDM2 expression.
BD138247
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Location/Qualifiers
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C12N15/00
Strandedness: Single,
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JP 2002508944-A/173.
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JP 2002508944-A/172.
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Best Local Similarity 100.
Matches 20; Conservative
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BD138247/c
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KEYWORDS
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KEYWORDS
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26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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CI2N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
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C12Q1/68,
C12N15/00
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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0.8%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
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Antisense modulation of human MDM2 expression.
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Location/Qualifiers
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7.9e+02;
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0.8%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 7.5
Matches 20; Conservative 0; Mismatches
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                                                 26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/04881
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Strandedness: Single;
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                 JP 2002508944-A/176
26-MAR-2002
   Unidentified
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BD138251/c
LOCUS
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ACCESSION
VERSION
KEYWORDS
SOURCE
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TITLE
JOURNAL
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1 (bases 1 to 20)

Mirablia, L. Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.

Antisense modulation of human MDM2 expression

Patent: JP 2002508944-A 175 26-MAR-2002;

ISIS PHARMAGUTICLES INC

S Unidentified

PN JP 2002508944-A/175

PD 36-MAR-1999

PP 26-MAR-1999

PP 26-MAR-1999

PP 26-MAR-1999

PP 26-MAR-1999

PP 26-MAR-1999

PP LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
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1 (bases 1 to 20)
Miraglla,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944.A 176 26-MAR-2002;
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Antisense modulation of human MDM2 expression FH
Location/Qualifier . 20
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                               Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 7.9e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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Antisense modulation of human MDM2 expression.
BD138249
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db_xref="taxon:32644"
                                                                                                  1439 GGAAATGATGATAAATTA 1458
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JP 2002508944-A/175.
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JP 2002508944-A/176.
unidentified
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DEFINITION ACCESSION VERSION KEYWORDS SOURCE

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AUTHORS TITLE JOURNAL

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DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM

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PAT 18-SEP-2002

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BD138254.1 GI:23233199

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Is 1 (bases 1 to 20)

Miragliat.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.

Antisense modulation of human MDM2 expression

Patent: JP 2002508944-A 180 26-MAR-2002;

ISIS PHARMACEUTIALS INC

OS Unidentified

PN JP 2002508944-A/180

PD 26-MAR-1999

PD 26-MAR-1999
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LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
                                                           CI2N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//C12Q1/68,
C12Q1/68,
C12N15/00
Strandedness: Single,
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C12D1/68,
C12N15/00
Strandedness: Single;
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Antisense modulation of human MDM2 expression FH Location/Qualifiers 1. 20
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Antisense modulation of human MDM2 expression FH Location/Qualifiers
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Antisense modulation of human MDM2 expression.
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Location/Qualifiers
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TITLE
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unclassified.

E 1 (bases 1 to 20)
S Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
L Patent: JP 2002508944-A 178 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/178
PD 26-MAR-1999 JP 2000538025
PF 26-MAR-1999 UP 2000538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
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LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
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unclassified.

I (bases 1 to 20)
Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 179 26-MAR-2002;
Sulidentified
PN JP 2002508944-A/179
PD 26-MAR-2002
PP 26-MAR-1999 JP 200538025
PR 26-MAR-1999 US 09/048810
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Antisense modulation of human MDM2 expression
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Antisense modulation of human MDM2 expression.
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Pred. No. 7.9e+02;
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V 100.0%; Pred. No. '..
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/db_xref="taxon:32644"
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JP 2002508944-A/178.
unidentified
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JP 2002508944-A/179.
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BD138252/c
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BD138258/c
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BD138257/c
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unclassified.

E 1 (bases 1 to 20)
S Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
L Patent: JP 2002508944-A 181 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/181
PD 26-MAR-2002
PF 26-MAR-199 JP 200538025
PR 26-MAR-199 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM, BRETT P MONIA,LEX M
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JP 200250844-A/182
26-MAR-1200
26-MAR-1999 JP 2000538025
26-MAR-1999 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
                                                                                                                                                                                                                                                      26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
                                       PAT 18-SEP-2002
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C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
C12N15/00
Strandedness: Single;
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
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Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 182 26-MAR-2002;
ISIS PHARMACEUTICALS INC
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
                             20 bp DNA linear Antisense modulation of human MDM2 expression.
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Antisense modulation of human MDM2 expression.
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Location/Qualifiers
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0.8%; Score 20; DB 1; L
Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;
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JP 200250894-A/182.
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JP 2002508944-A/181.
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BD138256/c
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AUTHORS
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Miragila, L. J., Nero, P., Graham, M. J., Monia, B. P. and Cowsert, L. M. Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 183 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2005508944-A/183
PD 26-MAR-2002
PP 26-MAR-1999 JP 200538025
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 10-NEN J MIRAGILA, PAMBLA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
Strandedness: Single,
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Strandedness: Single;
Topology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                           Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 7.9e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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Antisense modulation of human MDM2 expression.
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Antisense modulation of human MDM2 expression.
                                                                                                                                          /organism='Unidentified'.
Location/Qualifiers
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JP 2002508944-A/183.
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unclassified.

1 (bases 1 to 20)
Miraglia,L.J., Nerco,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense medulation of human MDM2 expression
Patent: JP 2002508944-A 186 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/186
PD 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 UP 2000538025
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 10CRU J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
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C12N15/00
Strandedness: Single;
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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                                                                                                                                   0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 7.9e+02;
tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                           20 bp DNA line
Antisense modulation of human MDM2 expression.
BD138260
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                                   Location/Qualifiers
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JP 2002508944-A/186.
unidentified
unidentified
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Best Local Similarity 100.0
Matches 20; Conservative
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BD138261/c
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BD138260/c
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                                            Unidentified

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I (bases 1 to 20)

S Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
L Patent: 19 2002508944-A 184 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/184
PD 26-MAR-1999 UP 2000538025
PR 26-MAR-1999 US 09/048810
PR 1 LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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JP 2002508944-A/185
26-MAR-2005
26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
                                                                                                                                                                                                                                26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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C12Q1/68,
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unclassified.

1 (bases 1 to 20)
Mixaglia-L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 185 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/185
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Strandedness: Single;
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               BD138258.1 GI:23233203
JP 2002508944-A/184.
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C12N15/00
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BD138259/c
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BD138263/c
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BD138264/c
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AUTHORS
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JOURNAL
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Antisense modulation of human MDM2 expression
Batent: JP 2002508944-A 187 26-MAR-2002;
S Unidentified
BN J2 2002508944-A/187
PD 26-MAR-2002
PP 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 10 PR J MIRAGILA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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                                                                                                                                 COWSERT
C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C1201/68,
C12N15/00
Strandednese: Single;
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C12Q1/68,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
                                                                                                                                                                                                                                                                                                            0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 7.9e+02; tive 0; Mismatches 0; Indels
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Strandedness: Single;
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Best Local Similarity 100.0%
Matches 20; Conservative
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ACCESSION
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KEYWORDS
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BD138262/c
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TITLE
JOURNAL
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AUTHORS
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BD13263.1 GI:23233208

BD13264.4/189.

Unidentified

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I (bases 1 to 20)

Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.

Antieense modulation of human MDM2 expression

L Patent: JP 2002508944-A 189 26-MAR-2002;

SS Unidentified

NOS Unidentified

PD 26-MAR-2002

PD 26-MAR-2002

PP 26-MAR-1999 JP 2000538025

PR 26-MAR-1999 US 099/048810

PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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C12Q1/68,
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JP 2002508944-A/190.
unidentified
unclassified.
1 (bases 1 to 20)
Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 190 26-MAR-2002;
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
                                                                     0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 7.9e+02;
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Antisense modulation of human MDM2 expression.
BD138263
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20; Conservative 0; Mismatches
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Strandedness: Single;
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DP 2002508944-A/192
26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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Cl2Q1/68,
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1 (bases 1 to 20)
Wiraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Miraglia,L.J., Nero,P., Graham, M.J., Monia,B.P. and Cowsert,L.M.
Patisense modulation of human MDM2 expression
Patent: JP 2002568944-A 193 26-MAR-2002;
Unidentified
N JP 2002568944-A/193
PP 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
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                                                                                                                           BD138266 20 bp DNA linear Antisense modulation of human MDM2 expression.
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     0; Indels
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Antisense modulation of human MDM2 expression.
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Location/Qualifiers
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                              1617 GAACCTTGTGTGATTTGTCA 1636
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                                                                                                                                                                                                                                                                                                                                                                                                                                              C12N15/00
Strandedness: Single;
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                                               GAACCTIGIGIGATITGICA 1
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JP 2002508944-A/193.
    20; Conservative
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     Matches
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AUTHORS
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TITLE
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FILE Antisense modulation of human MDM2 expression

FILE Patent: JP 2002508944-A 191 26-MAR-2002;

ENT OS Unidentified

PD 26-MAR-2002

PP 26-MAR-1999 JP ~

PR 26-MAR-1999 JP ~

PI 1
                                                  26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO,MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
Strandedness: Single,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Location/Qualifiers
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Location/Qualifiers
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Strandedness: Single;
                           JP 2002508944-A/190
26-MAR-2002
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  PHARMACEUTICALS INC
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BD138265/c
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AUTHORS
TITLE
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               COMMENT
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                            RESULT 649
BD138269/C
LOCUS
DEFINITION
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BD138270/c
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AUTHORS
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Unidentified
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I (bases 1 to 20)

Wixegliah.L.J. Nerco,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.

Antisense modulation of human MDM2 expression

L Patent: JP 2002508944-A 194 26-MAR-2002;

Unidentified
PN JP 2002508944-A/194
PN 2002508944-A/194
PN 2002508944-A/199
PN 2002508941-B99 UP 2000538025
PR 26-MAR-1999 UP 2000538025
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26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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C12Q1/68,
C12N15/00
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
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C12N15/00
Strandedness: Single,
Topology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
source 1. .20
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100.0%; Pred. No. 7.9e+02;
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                                                                                                                                                                    /organism='Unidentified'
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Location/Qualifiers
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Best Local Similarity 100.
Matches 20; Conservative
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BD138268/c
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BD138269.1 GI:3323314
BD138269.1 GI:3233214
JP 2002508944-A/195.
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Antisense modulation of human MDM2 expression
LD Patent: JP 2002508944-A 195 26-MAR-2002;
EM INVERSE INC
OS Unidentified
EMAR-2002
EMAR-2002
EMAR-1999 JP 2000538025
EMAR-1999 UP 2000538025
EMAR-1998 US 09/048810
ELOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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unidentified
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Later (Dases 1 to 20)

Miralla, L. J., Nero, P., Graham, M. J., Monia, B. P. and Cowsert, L. M.
Antisense modulation of human MDM2 expression

Later (JP 2002508944-A 196 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/196
PP 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 US 09/048810
PP 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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JP 2002508944-A/196
26-MAR-2002
26-MAR-1999 JP 200538025
26-MAR-1999 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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PAT 18-SEP-2002
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JP 2002508944-A/195
S26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1999 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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C12Q1/68,
C12N15/00
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C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
linear
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BD138269 20 bp DNA line
Antisense modulation of human MDM2 expression.
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7.9e+02;
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FEATURES

DEFINITION ACCESSION

ORGANISM

VERSION KEYWORDS SOURCE

REFERENCE AUTHORS TITLE JOURNAL

COMMENT

RESULT 651 BD138271/c LOCUS

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### 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 | 1913 
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Patent: JP 2002508944-A 199 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/199
PP 26-MAR-1999 UP 2000538025
PP 26-MAR-1999 UP 2000538025
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 10-REN J MIRAGILA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
C12N15/00
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C12NIS/00
Strandedness: Single;
Topology: Linear;
Antisense modulation of human MDM2 expression FH Key
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Antisense modulation of human MDM2 expression.
BD138273
          human MDM2 expression.
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/organism='Unidentified'
Location/Qualifiers
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/organism="unidentified"
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/db_xref="taxon:32644"
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JP 2002508944-A/199.
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BD138273/c
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26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
Strandedness: Single;
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Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2012508944-A 197 26-MAR-2002;
ISIS PHARNACEUTICALS INC
                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
BD138271
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/organism="unidentified"
/mol_type="genomic DNA"
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/db_xref="taxon:32644"

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C12Q1/68,
C12N15/00
Strandedness: Single;
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JP 2002508944-A/197.
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Best Local Similarity 100.0
Matches 20; Conservative
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1 (bases 1 to 20)

Miragliaci...J., Nercy, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.

Antisense modulation of human MDM2 expression

Patent: JP 2002508944-A 200 26-MAR-2002;

ISIS PHARMAGENTICALS INC

S Unidentified

PN JP 2002508944-A/200

PD 36-MAR-1999 JP 2000538025

PP 26-MAR-1999 US 09/048810

PP 26-MAR-1999 US 09/048810

PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
1. .20
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BD138275
BD138275. GI:23233220
JP 2002589544-A/201.
unidentified
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Antisense modulation of human MDM2 expression.
                 /organism='Unidentified'
Location/Qualifiers
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BD138274.1 GI:23233219
JP 2002508944-A/200.
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Best Local Similarity 100.0%
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1 (bases I to 20)
Miragila, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 202 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/202
PP 26-MAR-1999 JP 2000538025
PF 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
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unclassified.

1 (bases 1 to 20)
Miraglia, L.J., Nerro, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
Antisense modulation of human MDM2 expression
Batent: JP 2002508944-A 201 26-MAR-2002;
S. Unidentified
D. 2002508944-A/201
PD 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 JB 09/048810
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
C12N15/00
Strandedness: Single;
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
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Location/Qualifiers
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Strandedness: Single;
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JP 2002508944-A/202.
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BD138279/c
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JP 2002508944-A/203.

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E 1 (bases 1 to 20)

Antisense modulation of human MDM2 expression

L Patent: JP 2002508944-A 203 26-MAR-2002;

ISIS PHARMACEUTICALS INC

OS Unidentified

N JP 2002508944-A/203

PP 26-MAR-1999 JP 2000538025

PR 26-MAR-1999 US 099/048810

PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
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Antisense modulation of human MDM2 expression.
BD138278.1 GI:23233223
BD138278.2 GI:23233223
JP 2002508944-A/204.
unidentified
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unidentified.
I (bases 1 to 20)
Miraglia, L. J., Nero, P., Graham, M. J., Monia, B. P. and Cowsert, L. M.
Antisense modulation of human MDM2 expression
                                                                                      Gaps
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
                                                      Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 7.9e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                      linear
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Antisense modulation of human MDM2 expression.
BD138277
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Location/Qualifiers
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LOCUS
DEFINITION
ACCESSION
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BD138277/c
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TITLE
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Patent: JP 2002508944-A 204 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/204
PD 26-MAR-2002
PP 26-MAR-1999 JP 2000518025
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
C12N15/00
Strandedness: Single;
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C12D1/68,
C12N15/00
Strandedness: Single,
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Miraglia, L. J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
Pattisense modulation of human MDM2 expression
Battisense modulation of human MDM2 expression
OS Unidentified
OS Unidentified
PN JP 2002508444-A/205
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
BD138279
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Location/Qualifiers

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Gaps

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PAT 18-SEP-2002

linear

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Gaps

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Key

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uniclassified.

1 (bases 1 to 20)

Miraglia.L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 208 26-MAR-2002;
ISIS PHRAMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/208
PN JP 2002508944-A/208
PN 26-MAR-1999 UP 200538025
PR 26-MAR-1998 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
   26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
                                                                 COWSERT C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//C12N15/00 Strandedness: Single, Topology: Linear; Topology: Linear; Antisense modulation of human MDM2 expression FH Key Location/Qualifiers
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C12Q1/68,
C12N15/00
Strandedness: Single;
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Antisense modulation of human MDM2 expression.
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JP 2002508944-A/208.
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DEFINITION
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KEYWORDS
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BD138282/c
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AUTHORS
TITLE
JOURNAL
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BD138281.
BD 2002508944-A/207.
NISM unidentified
unclassified.
NRS Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
ALISE PHARMACEUTICALS INC
OS Unidentified

ND 2002508944-A 207 26-MAR-2002;
BD 26-MAR-2002

PD 26-MAR-2002
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26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
                                                                                                                                                                                                                                                                             unclassified.

1 (bases 1 to 20)
Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M. Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 206 26-MAR-2002;
                      Gaps
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Antisense modulation of human MDM2 expression Location/Qualifiers
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Antienne modulation of human MDM2 expression.
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Location/Qualifiers
 100.0%; Pred. No. 7.9e+02; ive 0; Mismatches 0;
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0.8%; Score 20; DB 1; L.
Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;

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JP 2002508944-A/206
26-MAR-2002
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JP 2002508944-A/206.
Best Local Similarity 100.
Matches 20; Conservative
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BD138280/c
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STATEMENTORITICALS

NATIONALY, NETO, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.

Antisense modulation of human MDM2 expression

Antisense modulation of human MDM2 expression

LESTS PHARMACEUTICALS INC

OS Unidentified

PN JP 2002508944-A/211

PD 26-MAR-2002

PP 36-MAR-1999 JP 2000538025

PR 26-MAR-1999 JP 2000538025

PR 26-MAR-1999 US 09/048810

PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
Strandedness: Single,
     C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
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                                                                                  Topology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
source 1. .20
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                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 20; DB 1; Length 20;
llarity 100.0%; Pred. No. 7.9e+02;
Conservative 0; Mismatches 0; Indels
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Antisense modulation of human MDM2 expression.
BD138285
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                           C1201/68,
C12N15/00
Strandedness: Single;
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Matches 20; Conserv
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26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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I (bases 1 to 20)
Miraglia, L. J., Nero, P., Graham, M.J., Monla, B.P. and Cowsert, L.M.
Antisense modularion of human MDM2 expression
Patent: JP 2002508944-A 209 26-MAR-2002;
SUNIGHOLIFIED INC
ON Unidentified
PN JP 2002508944-A/209
PD 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMBLA NERO, MARK J GRAHAM, BRETT P MONIA, LEX P
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                                                                                                                        linear
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                                                                                                                     BD138283 20 bp DNA line
Antisense modulation of human MDM2 expression.
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Location/Qualifiers

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JP 2002508944-A/209.
20 AGAATTATATATTCTAACT
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Best Local Similarity
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BD138284/c
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VERSION
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BD138288/c
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Patent: JP 200250844-A 213 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PD 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
C12N15/00
Strandedness: Single,
Topology: Linear;
Antisense modulation of human MDM2 expression FH Key
Location/Qualifiers
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C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C1201/69,
C12N15/00
Strandednese: Single,
                                                                                                      unclassified.

1 (bases 1 to 20)

Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 212 26-MAR-2002;
OS Unidentified
DS Unidentified
NP 2002508944-A/212
PD 26-WAR-1999 JP 200538025
PF 26-MAR-1999 US 09/048810
PF 26-WAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX
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20 bp DNA line
Antisense modulation of human MDM2 expression.
BD138286
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Location/Qualifiers

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                                               BD138286.1 GI:23233231
JP 2002508944-A/212.
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Best Local Similarity 100.0
Matches 20; Conservative
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BD138287/c
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C12Q1/68,
C12N15/00
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Antisense modulation of human MDM2 expression FH Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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BD138289.
BD138289.1 GI:23233234
JP 2002508944-A/215.
                                                                                                                                                             Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 7.9e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
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Location/Qualifiers
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Location/Qualifiers
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Antisense modulation of human MDM2 expression
Datent: JP 2002508944-A 217 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/217
PD 26-MAR-2002
PR 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
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                                                                                                       Gaps
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
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                                                                                                                                   1865 TATATCAAAGTGAAAATG 1884
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Strandedness: Single;
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JP 2002508944-A/218.
                                                                       Query Match
Best Local Similarity
Matches 20; Conserval
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BD138292/c
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BD138291/c
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unclassified.

E 1 (bases 1 to 20)
S Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
L Patent: JP 2002508944-A 216 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/216
PD 26-MAR-1999 JP 200538025
PR 26-MAR-1999 UF 200538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM, BRETT P MONIA, LEX M
                                                  Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M. Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 215 26-MAR-2002;
S Unidentified
PN JP 2002508944-A/215
PD 36-MAR-2002
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
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Antisense modulation of human MDM2 expression FH
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Antisense modulation of human MDM2 expression FH
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100.0%; Pred. No. 7.9e+02;
ative 0; Mismatches 0; Indels
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Location/Qualifiers

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Strandedness: Single;
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JP 2002508944-A/216.
unidentified
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Best Local Similarity 100.'
Matches 20; Conservative
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BD138290/c
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BD138294/c
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Antisense modulation of human MDM2 expression
Patent: JP 200268944A 218 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002568944-A/218
PD 36-MAR-2002
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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Patent: JP 2002508944-A 219 26-MAR-2002;
ISIS PHARMAGEUTICALS INC
S. Unidentified
BN JP 2002508944-A/219
BD 36-MAR-2002
BP 36-MAR-1999 JP 2000538025
BR 26-MAR-1999 US 09/048810
BI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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CIZN15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
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C12Q1/68,
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Antisense modulation of human MDM2 expression FH Location/Qualifiers
source 1. .20
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Antisense modulation of human MDM2 expression
Location/Qualifiers
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/organism="unidentified"
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Strandedness: Single;
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BD138293/c
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D138294.1 GI:21233239

D138294.1 GI:21233239

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I (bases 1 to 20)

Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression

Patent: JP 2002508944-A 220 26-MAR-2002;

Unidentified

D 26-MAR-2002

PN JP 2002508944-A/220

PP 26-MAR-1999 UP 2000538025

PR 26-MAR-1999 US 09/048810

PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
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C12Q1/68,
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Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 221 26-MAR-2002;
ISIS PHARNACEUTICALS INC
OS Unidentified
PN JP 2002
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 7.9e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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BD138295.1 GI:23233240
JP 202058944-A/221.
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Antisense modulation of human MDM2 expression.
BD138294
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Location/Qualifiers
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7.9e+02;
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                                                                              1889 AATTCACATAGATTTCTTCT 1908
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BD138298/c
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LOCUS
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TITLE
JOURNAL
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TITLE
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BD138296.1 GI:22233241
JP 2002508944-A/222.
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IE (bases 1 to 20)

Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Antisense modulation of human MDM2 expression

ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/222
PN 26-MAR-2002
PN 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO,MARK J GRAHAM, BRETT P MONIA,LEX M
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C12Q1/68,
C12N15/00
                                                                              C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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llarity 100.0%; Pred. No. 7.9e+02;
Conservative 0; Mismatches 0; Indels
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Antisense modulation of human MDM2 expression.
BD138296
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Location/Qualifiers

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Strandedness: Single;
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Best Local Similarity
Matches 20; Conserv
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BD138297.1 GI:23233242

BD138297.1 GI:23233242

JP 2002508944-A/223.

Windentified

Unclassified.

SM iraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M..

Antisense modulation of human MDM2 expression

AL Patent: JP 2002508944-A 223 26-MAR-2002;

ISIS PHARMACEUTICALS INC

OS Unidentified

PN JP 2002508944-A/223

PD 26-MAR-1999 JP 2000538025

PR 26-MAR-1999 US 09/048810

PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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SSION BD138298.

BD138298.1 GI:23233243

NDS JP 2002508944-A/224.

"B unidentified unclassified.

"ANISM unidentified unclassified.

"A D 2002508944-A 224 26-MAR-2002;

"B Antisense modulation of human MDM2 expression

"A D 2002508944-A/224

"B D 26-MAR-1999 JP 2000538025

"B C 26-MAR-1999 JP 2000538025
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26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12N1S/09, A61K48/00, A61P9/10, A61P17/06, A61P3S/00, C07H21/04//
C12Q1/68,
C12N1S/00
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Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
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Location/Qualifiers
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Antisense modulation of human MDM2 expression
Batent: JP 2002508944-A 225 26-MAR-2002;
Sunidentified
BN JP 2002508944-A/225
PP 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 UP 2000538025
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGILA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
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                                                  Strandedness: Single;
Topology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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100.0%; Pred. No. 7.9e+02;
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Antieense modulation of human MDM2 expression.
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Location/Qualifiers
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JP 2002508944-A/225.
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RESULT 680

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Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M. Antiaglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M. Antiagnae modulation of human MDM2 expression
Patent: JP 2002508944-A 227 26-MAR-2002;
S Dindentified
S Unidentified
DP 26-MAR-2002
PP 26-MAR-1999 JP 200538025
PF 26-MAR-1999 US 099/048810
PR 26-MAR-1999 US 099/048810
PR 10-REN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
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           PAT 18-SEP-2002
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                                                                                                                                    unclassified.

1 (bases 1 to 20)

Miraglia, L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.

Antisense modulation of human MDM2 expression

Patent: JP 2002508944-A 226 26-MAR-2002;

ISIS PHARMACETUTCALS INC

ON Unidentified

PR 2002508944-A/226

PP 26-MAR-1999 JP 2000538025

PP 26-MAR-1999 US 09/048810

PR 26-MAR-1999 US 09/048810

PR 26-MAR-1999 US 09/048810

PR 10-OREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX I
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C12Q1/68,
C12N15/00
Strandedness: Single;
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
C12N15/00
Strandedness: Single;
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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20 bp DNA line
Antisense modulation of human MDM2 expression.
BD138300
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Antisense modulation of human MDM2 expression.
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Location/Qualifiers
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0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;
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JP 2002508944-A/226.
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JP 2002508944-A/227.
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BD138301/c
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JOURNAL
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AUTHORS
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JOURNAL
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PEATURES

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Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M. Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 229 26-MAR-2002;
Olidentified
PN JP 2002508944-A/229
PN JP 2002508944-A/229
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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1 (bases 1 to 20)

Miraplia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.

Antisense modulation of human MDM2 expression

Patent: JP 2002508944-A 230 26-MAR-2002;

ISIS PHARMAGEUTICALS INC

S Unidentified

PR 2002508944-A/230

PP 26-MAR-1999 UP 2000538025

PP 26-MAR-1999 US 09/048810

PR 26-MAR-1999 US 09/048810

PR 26-MAR-1999 US 09/048810

PR 10REN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
                                                                                                                                                                                                                                                                                                                                                                                      COWSERT
C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
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C12D1/68,
C12N15/00
Strandedness: Single,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
BD138304
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C12N15/00
Strandedness: Single;
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JP 2002508944-A/230.
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       JP 2002508944-A/229
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1 (bases 1 to 20)
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BD138304/c
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AUTHORS
TITLE
JOURNAL
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       KEYWORDS
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BB138302 1 G1:23233247

BD138302 1 G1:23233247

JP 2002508944-A/228.

unidentified unclassified.

I (bases 1 to 20)

Antisense modulation of human MDM2 expression

L (bases 1 to 20)

Antisense modulation of human MDM2 expression

L STRARMACEUTICALS INC

OS Unidentified

DP 2002508944-A/228

PN JP 2002508944-A/228

PN JP
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26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
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Topology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
BD138303
BD138303.1 GI:23233248
                                                                                             /organism='Unidentified'.
Location/Qualifiers
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Location/Qualifiers
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100.0%; Pred. No. 7.9e+02;
ative 0; Mismatches 0;
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/db_xref="taxon:32644"

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    /organism="unidentified"
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/db_xref="taxon:32644"

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BD138302/c
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TITLE JOURNAL AUTHORS REFERENCE

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BD138307/c
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                 TITLE
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1P 200250844-A/231
26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1999 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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CI2Q1/68,
CI2NI5/00
                                                                                                                                                                                                                                                                                                              unidentified
unclassified.
1 (bases 1 to 20)
Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 231 26-MAR-2002;
ISIS PHARMACEUTICALS INC
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                                                                               Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 7.9e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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BD138306.1 GI:23233251
JP 2002508944-A/232. unidentified undentified undentified undentified undentified.
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Antisense modulation of human MDM2 expression.
BD138305
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0.8%; Score 20; DB 1; Lv
Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;

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    /mol_type="genomic DNA"
    /db_xref="taxon:32644"

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/mol_type="genomic DNA"
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  location/Qualifiers
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JP 2002508944-A/231.
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BD138305/c
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JP 2002508944-A/233
26-WAR-1200
26-WAR-1999 JP 2000538025
26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 232 26-MAR-2002;
ISIS PHARMAGEUTICALS INC
OS Unidentified
PN JP 2002508944-A/232
PD 36-MAR-2002
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1998 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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C12Q1/68,
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
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Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Partisense modulation of human MDM2 expression
Patent: JP 2002508944-A 233 26-MAR-2002;
ISIS PHARMACEUTICALS INC
S Unidentified
PN JP 2002508944-A/233
PD 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM, BRETT P MONIA.LEX
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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BD138307
BD138307.1 GI:23233252
JP 200250944-A/233.
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Location/Qualifiers
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/mol_type="genomic DNA"
/db_xref="taxon:32644"
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/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"
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Strandedness: Single;
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Strandedness: Single;
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26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
JP 2002508944-A/235
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JP 2002508944-A/236.
unidentified
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Best Local Similarity 100.
Matches 20; Conservative
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BD138310/c
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S Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
L Patent: JP 2002208944-A, 234 26-MAR-2002;
ISIS PHARNACETTICALS INC
O Unidentified
PN JP 2002508944-A/234
PD 26-MAR-1999 UP 2000538025
PF 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
Strandedness: Single;
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1 (bases 1 to 20)
Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Parent: JP 2002508944-A 235 26-MAR-2002;
ISIS PHARMAGENTICALS INC
OS Unidentified
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Antisense modulation of human MDM2 expression
          Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 7.9e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                  linear
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BD138309
BD138309.1 GI:23233254
DIP 2002508944-A/235.
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Antisense modulation of human MDM2 expression.
BD138308
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LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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Patent: JP 2002508944-A 236 26-MAR-2002;
S Unidentified DN DR 2002508944-A/236
PD 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX P
                                                           C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
C12N15/00
Strandedness: Single,
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C12D1/68,
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Strandedness: Single,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
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100.0%; Pred. No. 7.9e+02;
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BD138313/c
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LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C1201/68,
C12N15/00
                                                                                                                                        unclassified.
(Description of Mana, M.J., Monia, B.P. and Cowsert, L.M. Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M. Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 237 26-MAR-2002;
ISIS PHARMACEUTICALS INC
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Antisense modulation of human MDM2 expression
Location/Qualifiers
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Antiennee modulation of human MDM2 expression.
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JP 2002508944-A/237
26-MAR-1999 JP 2000538025
26-MAR-1999 US 09/048810
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          20 TCTGTCTTAAATGAGAAGTA 1
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JP 2002508944-A/237.
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1 (bases 1 to 20)
Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 239 26-MAR-2002;
ISIS PHARMAGENTICALS INC
BY Unidentified
BY JP 2002508944-A/239
BP 26-MAR-1999 JP 2000538025
BP 26-MAR-1999 UP 200538025
BP 26-MAR-1999 US 09/048810
BR 26-MAR-1999 US 09/048810
BI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX P
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C12D1/68,
C12N15/00
Strandedness: Single,
C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
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Strandedness: Single,
Strandedness: Linear;
Topology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression
L Patent: UP 20022508944-A 342 26-MAR-2002;
ISIS PHARMACETTICALS INC
OS Unidentified
PN JP 2002508944-A/242
PD 26-MAR-1999 UP 2000538025
PF 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BREȚT P MONIA, LEX M
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LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BREȚT P MONIA, LEX M
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C12Q1/68,
                      Topology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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BD138316 BD13816.1 GI:23233261
JP 2002508944-A/242.
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100.0%; Pred. No. 7.9e+02;
tive 0; Mismatches 0;

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AUTHORS Miraglia, LJ. J. Nero, P. Graham, M.J., Monia, B.P. and Cowsert, L.M. FILE Antisense modulation of human MDM2 expression
FITLE Antisense modulation of human MDM2 expression
FITLE Patent: JP 2002508944-A 241 26-MAR-2002;
SNT OS Unidentified
PN JP 2002508944-A/241
PD 26-MAR-2002
PP 26-MAR-1999 JP JP
PR 26-MAR-1999 JP JP
PR 26-MAR-1999 JP JP
PR 26-MAR-1002
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26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, WARK J GRAHAM, BRETT P MONIA, LEX M
                                                                                                                                                                                                                                                                             26-MAR-1999 UP 2000538025
26-MAR-1999 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
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C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
C12N15/00
                                                                                                                                                                    Miragila,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M. Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 240 26-MAR-2002;
ISIS PHARACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/240
PD 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1998 US 09/048810
PI LOREN J MIRAGILA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA.IRX
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Strandedness: Single;
Topology: Linear;
Antisense modulation of human MDMZ expression FH
Location/Qualifiers
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Pred. No. 7.9e+02;
                                   linear
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                                 BD138314 20 bp DNA lin
Antisense modulation of human MDM2 expression.
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Best Local Similarity 100.0%; Pred. No. 7.9
Matches 20; Conservative 0; Mismatches

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/db_xref="taxon:32644"

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BD138314.1 GI:23233259
JP 2002508944-A/240.
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BD138315/c
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Strandedness: Single;
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JP 2002508944-A/245.
unidentified
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JP 2002508944-A/246.
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unclassified.
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BD138319/c
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JOURNAL
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                                                                                                        Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Autisense modulation of human MDM2 expression
Patent: JP 2002508944-A 243 26-WAR-2002;
SS Unidentified
NJ 2002508944-A/243
PD 26-WAR-2002
PP 26-WAR-2002
PP 26-WAR-1999 JP 2000538025
PR 26-WAR-1999 US 09/048810
PR 26-WAR-1999 US 09/048810
PR LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX M
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TITLE Antisense modulation of human MDM2 expression
FOURNAL ISS PHARMACEUTICALS INC
ENT OS Unidentified
PD 26-MAR-2002
PP 26-MAR-1999 JP --
PR 2
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26-WAR-1903
26-WAR-1999 JP 2000538025
26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
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C12Q1/68,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Pred. No. 7.9e+02;
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BD138318.1 GI:23233263
JP 20025089444-2/72
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Strandedness: Single;
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Strandedness: Single;
    BD138317.1 GI:23233262
JP 2002508944-A/243.
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unclassified.
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S Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
I Patent: JP 2002508944-A 245 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/245
PP 26-MAR-1999 JP 2005538025
PP 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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JP 2002508944-A/245
26-MAR-2002
26-MAR-1999 JP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
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7.7.9e+02; o; Indels
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Antisense modulation of human MDM2 expression.
BD138319
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Antisense modulation of human MDM2 expression.
BD138320
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Location/Qualifiers
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PAT 18-SEP-2002

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26-WAR-1999 JP 2000538025
26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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C12Q1/68,
C12N15/00
Strandedness: Single,
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Up 2002508944-A/249.

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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
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/organism='Unidentified'
Location/Qualifiers
                                                        Query Match 0.8%; Score 20; DB 1; L. Best Local Similarity 100.0%; Pred. No. 7.9e+02; Matches 20; Conservative 0; Mismatches 0;
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Matches 20; Conserv
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BD138322/c
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BD138321.1 GI:22233266

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Is bases 1 to 20)

Antisense modulation of human MDM2 expression

Antisense modulation of human MDM2 expression

LISS PHARMACEUTICALS INC

OS Unidentified

PN JP 2002508944-A/247

PN 26-MAR-1999 JP 2000538025

PR 26-MAR-1999 US 09/048810

PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
                                                                                                                                     JP 2002508944-A/246
26-WAR-2002
26-WAR-1999 JP 2000538025
26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
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Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M. Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 246 26-MAR-2002;
SS Unidentified
PN JP 2002508944-A/246
PN JP 2002508944-A/246
PN JP 2002508944-A/246
PN JP 200250894-A/246
PN 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 095/048810
PR 26-MAR-1999 US 095/048810
PR 26-MAR-1999 US 095/048810
PR 26-MAR-1999 US 095/048810
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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100.0%; Pred. No. 7.9e+02;
ative 0; Mismatches 0; Indels
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Antisense modulation of human MDM2 expression.
BD138321
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Best Local Similarity
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BD138326/c
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JP 2002508944-A/250
26-MAR-1200
26-MAR-1999 JP 2000538025
26-MAR-1999 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
Unidentified
JP 2002508944-A/249
26-MAR-2002,
26-MAR-1999 UP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
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1 (Dasss 1 to 20)

Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 250 26-MAR-2002;
ISIS PHARMACEUTICALS INC
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
BD138324
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Location/Qualifiers

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JP 2002508944-A/250.
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C12N15/00
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AUTHORS
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BD138325

ACCESSION Antisense modulation of human MDM2 expression.

BD138325.

VERSION BD138325.1 GI:23233270

VERSION BD138325.1 GI:23233270

KEYWORDS Unidentified
Unidentified
Unclassified.

REFERENCE (CONSORDIAL)

AUTHORS ALTISE Antisense modulation of human MDM2 expression

TITLE Antisense modulation of human MDM2 expression

SOURCE ALTICE Antisense Modulation of human MDM2 expression

FITLE Antisense Modulation of human MDM2 expression

SOURCE ACCESSOR ANTISENSE ANTI
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JP 2002508944-A/251
26-MAR-2002
26-MAR-1999 UP 2000538025
26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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1 (bases 1 to 20)
Miraglia L.J., Mero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 252 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/252
PD 26-MAR-2002
PP 26-MAR-1999 JP 200538025
PR 26-MAR-1999 US 09/048810
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C12Q1/68,
C12N15/00
Strandedness: Single;
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Antisense modulation of human MDM2 expression.
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Location/Qualifiers
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/organism="unidentified"
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JP 2002508944-A/252.
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FEATURES

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PAT 18-SEP-2002

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1 (bases 1 to 20)
Miraglia, L.J., Nerv.P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 254 26-MAR-2002;
ISIS PHARM-EUTICALS INC

BY J 2002508944-A/254

PD 26-MAR-1999 JP 2000538025

PR 26-MAR-1999 US 09/048810

PR 26-MAR-1999 US 09/048810

PR 10-REN J MIRAGLIA, PAMMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COWSERT
C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
C12N15/00
Strandedness: Single;
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CI2N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Topology: Linear;
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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                                                Antisense modulation of human MDM2 expression.
BD138128 BD138128 BD138128 I GI:23233273
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Antisense modulation of human MDM2 expression.
BD138329
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BD138328/c
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BD138329/c
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SM unidentified
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I (bases i to 20)
SM miraglia.L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
L Patent: JP 2002508944-A 253 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/253
PD 26-MAR-2002
PP 26-MAR-2002
PR 26-MAR-1999 UP 200538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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JP 2002508944-A/253
JP 2002508944-A/253
26-MAR-1909
26-MAR-1999 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P. MONIA, LEX
                                        COWSERT
C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
C12N15/00
Strandedness: Single,
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C12Q1/68,
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Antisense modulation of human MDM2 expression
Location/Qualifiers
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Antisense modulation of human MDM2 expression
Location/Qualifiers
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Antieense modulation of human MDM2 expression.
BD138127 GI:23233272
JP 2002508944-A/253.
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Pred. No. 7.9e+02;
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/db_xref="taxon:32644"

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/db_xref="taxon:32644"

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BD138332/c
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26-MAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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CI2N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//
C12Q1/68,
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JP 2002508944-A/256.
unidentified
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l bases 1 to 20)
Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 256 26-MAR-2002;
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                         Topology: Linear,
Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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LOCUS BD138331 20 bp DNA line
DEFINITION Antisense modulation of human MDM2 expression.
                                                                          /organism='Unidentified'.
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Antiense modulation of human MDM2 expression.
BD138330
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Location/Qualifiers
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0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;
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/organism="unidentified"

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C12N15/00
Strandedness: Single;
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JP 2002508944-A/256
26-MAR-2002
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I (bases 1 to 20)

Mragliat, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.

Antisense modulation of human MDM2 expression

Patent: JP 2002508944-A 257 26-MAR-2002;

Unidentified

N JP 2002508944-A/257

PN JP 2002508944-A/257

PP 26-MAR-1999 JP 2000538025

PR 26-MAR-1999 US 09/048810

PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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Patent: JP 2002208944-A 28 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/258
PD 26-MAR-1999 JP 200538025
PP 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 10 CREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
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C12Q1/68,
C12NIS/00
Strandedness: Single;
Strandedness: Linear;
Antisense modulation of human MDM2 expression FH Key
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
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Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;
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Strandedness: Single;
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BD138332.1 GI:23233277
JP 2002508944-A/258.
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BD13833.1 GT:23233278

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Antisense modulation of human MDM2 expression
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100.0%; Pred. No. 7.9e+02;
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Antisense modulation of human MDM2 expression.
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/organism='Unidentified'
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0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;
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DNA linear PAT 18-SEP-2002

Antisense modulation of human MDM2 expression.

BD138335.1 G1:23233280

BD138335.1 G1:23233280

JP 2002508944-A/261.

Unidentified

unclassified.

LDasss 1 to 20)

Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.

Antisense modulation of human MDM2 expression

D Patent: JP 2002508944-A/261

LOSS Unidentified

PN JP 2002508944-A/261

PN JP 
Miragila, L. J. Nero, P. Graham, M. J., Monia, B. P. and Cowsert, L. M. Antisqila, L. J., Nero, P. Graham, M. J., Monia, B. P. and Cowsert, L. M. Antisquase modulation of human MDM2 expression
Patent: JP 2002508944-A 260 26-MAR-2002;
S Unidentified
PN J 2002508944-A/260
PP 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 UP 2000538025
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGILA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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C12Q1/68,
C12N15/00
Strandedness: Single;
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C12Q1/68,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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100.0%; Pred. No. 7.9e+02;
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BD138338/c
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ON Antisense modulation of human MDM2 expression.

BD138336 GT:23233281

JP 2002508944-A/262.

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MATISENSE modulation of human MDM2 expression

L Patent: JP 2002508944-A 262 26-MAR-2002;

SO Unidentified

D 26-WAR-2002

PD 26-WAR-2002

PD 26-WAR-1999 JP 2000538025

PR 26-WAR-1999 US 09/048810

PI LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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CI2N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//
C12Q1/68,
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Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 263 26-MAR-2002;
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Topology: Linear;
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Location/Qualifiers
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BD138337
BD188377.1 GI:23233282
JP 2002508944-A/263.
                                    ch 0.8%; Score 20; DB 1; Length 20; 1 Similarity 100.0%; Pred. No. 7.9e+02; 20; Conservative 0; Mismatches 0; Indels
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1 (bases 1 to 20)
Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 264 26-MAR-2002;
ISIS PHARMACEUTICALS INC
ON Unidentified
PN JP 2002508944-A/264
PD 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX I
                                                           26-MAR-1999 JP 2000538025
26-MAR-1999 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
                                                                                                                                                                       COWSERT C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//C12N15/09, A61K48/00, A61P9/10, A61P17/06, A61P35/00, C07H21/04//C12N15/00 Strandedness: Single, Topology: Linear; Antisense modulation of human MDM2 expression FH Key Location/Qualifiers
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/organism="unidentified"
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Strandedness: Single,
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BD138338.1 GI:23233283
JP 2002508944-A/264.
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                   Unidentified
JP 2002508944-A/263
PHARMACEUTICALS INC
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Matches

ઠ 셤 BD138339/c LOCUS

DEFINITION

ACCESSION VERSION

ORGANISM

KEYWORDS SOURCE

REFERENCE AUTHORS TITLE JOURNAL

COMMENT

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Miraglia, L. J., Nero, P., Graham, M. J., Monia, B. P. and Cowsert, L. M. Antisaglia, L. J., Nero, P., Graham, M. J., Monia, B. P. and Cowsert, L. M. Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 267 26-MAR-2002;
S Unidentified
BN JP 2002508944-A/267
PD 26-MAR-1999 19 2000538025
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PR 16-NEN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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 26-MAR-1998 US 09/048810-
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
                                                                        C12N15/09,A61K48/00,A61P9/10,A61P17/06,A61P35/00,C07H21/04//C12Q1/68,
C12Q1/68,
C12N15/00
Strandedness: Single,
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C12Q1/68,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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JP 2002508944-A/267.
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/organism="unidentified"
/mol_type="genomic DNA"
/db_xref="taxon:32644"

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    /organism="unidentified"
/mol_type="genomic DNA"
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BD138341/c
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26-MAR-1999 JP 2000538025
26-WAR-1998 US 09/048810
LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX
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C12Q1/68,
                                                                                                                                                                                                                                                                                                                  Miragila, L.U. 20)

Miragila, L.U. Wero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944-A 265 26-MAR-2002;
SUINGENTIFIED TO COMPANACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/265
PN JP 2002508949-A/265
PP 26-MAR-1999 JP 2000538025
PP 26-MAR-1999 UP 2000538029
PR 26-MAR-1999 UP 09/048810
PI LOREN J MIRAGILA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA. L.RX
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Miraglia, L.J., Nero, P., Graham, M.J., Monia, B.P. and Cowsert, L.M.
Antisense modulation of human MDM2 expression
Patent: JP 2002508944 266 26-MAR-2002;
ISIS PHARMACEUTICALS INC
OS Unidentified
PN JP 2002508944-A/266
PN JP 2002508944-A/266
PP 26-MAR-1999 JP 2000538025
   Gaps
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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Antisense modulation of human MDM2 expression.
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Antisense modulation of human MDM2 expression.
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Location/Qualifiers
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Pred. No. 7.9e+02;
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 0; Mismatches

    .20
    /organism="unidentified"
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                                    2319 TGATCCGCCCACCTCGGCCT 2338
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Strandedness: Single;
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JP 2002508944-A/266.
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Bource

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ACCESSION VERSION KEYWORDS SOURCE ORGANISM

REFERENCE AUTHORS TITLE JOURNAL

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Patent: JP 2002508944-A 268 26-MAR-2002;
S Unidentified
OS Unidentified
PD 26-MAR-2002
PP 26-MAR-1999 JP 2000538025
PR 26-MAR-1999 US 09/048810
PR 26-MAR-1999 US 09/048810
PI LOREN J MIRAGLIA,PAMELA NERO,MARK J GRAHAM,BRETT P MONIA,LEX N
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C1ZQ1/68,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chen, Y. Z., Hayashi, Y., Wu, J. G., Takaoka, E., Maekawa, K., Watanabe, N., Inazawa, J., Hosoda, F., Arai, Y., Mizushima, H., Morohashi, A., Ohira, M., Nakagawara, A., Liu, S., Hoshi, M., Horii, A. and Soeda, E.
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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                                                             linear
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N BD138342
BD138342
TP 200250894-A/268.
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/organism='Unidentified'
Location/Qualifiers
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/organism="unidentified"
/mol_type="genomic DNA"
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Genomics 74 (1), 55-70 (2001)
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Direct Submission
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Submitted (04-AUG-2001) Akira Horii, Tohoku University School of Medicine, Molecular Pathology; 2-1 Seiryomachi, Aoba-ku, Sendai, Miyagi 980-8575, Japan (E-mail:horii@mail.cc.tohoku.ac.jp, Tel:81-22-717-8042, Fax:81-22-717-8047)
Location/Qualifiers
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/organism='Unidentified'.
Location/Qualifiers
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C12N15/09,C12Q1/68,C12Q1/68,C12N15/00
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0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 7.9e+02;
Matches 20; Conservative 0; Mismatches 0;
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Best Local Similarity 100.0%; Pred. No. 8.2e+02;
Matches 20; Conservative 0; Mismatches 0;
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Sequence 27 from Patent WO0159152.
AX214484 AX214484.1 GI:15524532
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                                                       Zanger,U.M. and Lang,T.
Polymorphisms in the human cyp2b6 gene and their use in diagnostic and therapeutic applications
Patent: WO 0159152-A 27 16-AUG-2001;
Epidauros Biotechnologie AG (DE)
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
mdz12
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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/note="artificial sequence"
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S Agrawal, S. and Kandimalla, B.R.
Pseudo-cyclic oligonucleobases

AL Patent: US 6383752-A 12 07-MAY-2002;

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Sequence 5747 from Patent EP1281758.
AX693015
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Sequence 12 from patent US 6383752.
AR208396.1 GI:21509539
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synthetic construct
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Homo sapiens
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ch 0.8%; Score 20; DB 1; Length 26; ll Similarity 95.2%; Pred. No. 8.6e+02; 20; Conservative 0; Mismatches 1; Indels
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Best Local Similarity 91.3%; Pred. No. 8.4e+02;
Matches 21; Conservative 0; Mismatches 2;
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Method
Patent: WO 2004005547-A 138 15-JAN-2004;
IMPERIAL COLLEGE INNOVATIONS LIMITED (GB)
Location/Qualifiers
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/db xref="texon.32630"
/note="HS consensus sequence"
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Sequence 138 from Patent WO2004005547.
CQ766177
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Cullen, P. and Seedorf, U.
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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Sherr,C.J., Quelle,D., Roussel,M.F., Zindy,F. and Weber,J.D. ARF-P19, a novel regulator of the mammalian cell cycle
Patent: US 6407062-A 26 18-JUN-2002;
Location/Qualifiers
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Mutation of rad51 gene and its use in the diagnosis of predisposition to breast cancer
Patent: WO 0118254-A 13 15-MAR-2001;
THE DEPARTMENT OF HEALTH & HUMAN SERVICES (US)
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                                    Sequence 26 from patent US 6407062.
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Sequence 13 from Patent WO0118254.
AX093775
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AX612650.1 GI:28408079
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and mdz12
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Best Local Similarity 91.3%; Pred. No. 8.7e+02;
Matches 21; Conservative 0; Mismatches 2;
Coronary chip
Patent: WO 02072882-A 3675 19-SEP-2002;
OGHAM GmbH (DE)
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Sequence 5566 from Patent EP1281758.
AX692834 AX692834.1 GI:29415797

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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Genetic test to determine non-responsiveness to statin drug
treatment
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Genetic test to determine non-responsiveness to statin drug
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                            26 bp D Sequence 33 from patent US 6297014. AR171124
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Location/Qualifiers
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Sequence 33 from Patent EP1408121.
CQ798598.1 GI:46426960

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/organism="unknown"
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and mdz12
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Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
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Query Match 0.8%; Score 19.8; DB 1; Length 25; Best Local Similarity 91.3%; Pred. No. 8.7e+02; Matches 21; Conservative 0; Mismatches 2; Indels
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Sequence 5721 from Patent EP1281758.
AX692989
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Sequence 5648 from Patent EP1281758.
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Burgesb, C.E., Conley, P.B., Grosse, W.M., Hart, M., Kekuda, R., Shimkets, R.A., Spytek, K.A., Szekeres, B.S., Tomlinson, J.E., Topper, J.N. and Yang, R.B.
Proteins and nucleic acids encoding same Patent: WO 0216599-A 111 28-FEB-2002;
Curagen Corporation (US); COR THERAPEUTICS, INC. (US)
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/note="oligonuclectide primer"
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     0; Mismatches
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Patent: US 6228345-A 1 08-MAY-2001;
Location/Qualifiers
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Sequence 61 from patent US 6337192.
AR182144
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/organism="unknown"
/wol_type≈"unassigned DNA"
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Sequence 1 from patent US 6228345.
AR148944
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                                      2269 AGACAGGGTTTCACCGTGTTAGCCAG 2294
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                                                                             26 ACATGGGGTCTCACCGTGTTAGCCAG
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synthetic construct
artificial sequences.
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22; Conservative
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1 (bases 1 to 21)
   22; Conservative
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Matches 22; Conserv
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AR148944/c
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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Genetic test to determine non-responsiveness to statin drug
treatment
Weissman, S.M. and Jonsson, J.J.
Allele frequency differences method for phenotype cloning
Patent: US 650562-A 7 14-JAN-2003;
Location/Qualifiers
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Novel polypeptides and nucleic acids encoding same
Patent: WO 0144287-A 55 21-JUN-2001;
Curagen Corporation (US)
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/mol_type="unassigned DNA"
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/note="2826468 expression probe primer"
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84.6%; Pred. No. 8.9e+02;
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Defent: WO 0102606-A 33 11-JAN-2001;
Cedars-Sinai Medical Center (US)
Location/Qualifiers

    .26
    /organism="Homo sapiens"
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/db_xref="taxon:9606"

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Sequence 33 from Patent WO0102606.
AX068482
AX068482.1 GI:12578607
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Sequence 55 from Patent WO0144287.
AX190637
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                                                                       1. .26
/organism="unknown"
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RESULT 741 AX190637/c LOCUS DEFINITION

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RESULT 740 AX068482/c

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KEYWORDS SOURCE ORGANISM

REFERENCE AUTHORS TITLE

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PAT 20-APR-2002

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AX741033/c
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AX676183
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Genotyping reagents, kits and methods of use thereof Patent: WO 0129262-A 2866 26-APR-2001, Orchid BioSciences, Inc. (US)
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/organism="synthetic construct"
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                                                 Unclassified.

1 (bases 1 to 21)
Bartel, P.L. and Tavtigian, S.V.
MWSC1-an MWAC1 interacting protein
Patent: US 6337192-A 61 08-UAN-2002;
                                                                                                                                                                                                                                                                                                                                        Sequence 1318 from Patent WO0129262.
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Sequence 2866 from Patent WO0129262.
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/wol_type="unassigned DNA"
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/db_xref="taxon:32630"

    .21
    /organism="synthetic const:
/mol_type="unassigned DNA"

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 AR182144.1 GI:20225060
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Matches 20; Conservative
                          Unknown.
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AX116195
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Yan,W.L.
A method for producing a population of homozygous stem cells having a pre-selected immunophenotype and/or genotype
Patent: WO 0205429-A 40 25-JUL-2002;
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// Organism="synthetic construct"
// nol_type="genomic DNA"
// Mol_type="genomic DNA"
// Mol_type="genomic DNA"
// Mol_type="genomic DNA"
// NoTe="Description of Combined DNA/RNA Molecule:Synthetic
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Methods, kits and compositions pertaining to the suppression of
detectable probe binding to randomly distributed repeat sequences
in genomic nucleic acid
Patent: WO 03027328-A 7 03-APR-2003;
Boston Probes, Inc. (US); DakoCytomation Denmark A/S (DK)
Location/Qualifiers
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Pred. No. 8.5e+02;
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1 Similarity 95.2%; Pred. No. 8.5e+02;
20; Conservative 0; Mismatches 1;
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Sequence 40 from Patent WO02057429.
AX676183
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Sequence 7 from Patent WO03027328.
AX741033
/db_xref="taxon:32630"
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Location/Qualifiers
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il Similarity 95.2%;
20; Conservative
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artificial sequences.

I (bases 1 to 21)

S Lifton, R. P. and Simon, D. B.

Method to diagnose and treat pathological conditions resulting from deficient ion transport

L Patent: JP 2001508291-A 38 26-JUN-2001;

YALE UNIVERSITY

OS Artificial Sequence
BN JP 2001508291-A/38

PF 19-DEC-1997 JP 1998530123

PR 31-DEC-1996 US 08/778052

PI RICHARD P LIFTON, DAVID B SIMON

PC CIRNIS/09, CO7K14/435, CO7K16/00, C12N1/15, C12N1/19, C12N1/21, PC
BD056581 21-AUG-2002 Method to diagnose and treat pathological conditions resulting from deficient ion transport.
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synthetic construct
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attificial sequences.

1 (bases 1 to 22)
Mineno,J., Asada,K., Kato,I., Tanabe,C., Sasaki,H. and Terada,M.
Method for amplifying DNA
Patent: JP 2002345466-A 50 03-DEC-2002;
TAKARA BIO INC,THE PRESIDENT OF NATIONAL CANCER CENTER JAPAN, THI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Attificial Sequence
JP 2002345466-A/50
03-DEC-2002
08-MAY-2001 JP 2001137858
JUNICHI MINENO, KIYOZO ASADA, IKUNOSHIN KATO, CHIKAKO TANABE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PC C12221/02,C12Q1/68,G01N33/53,C12N15/00,C12N5/00 CC for analysis of human TSC gene Location/Qualifiers.

Location/Qualifiers
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1. .22
/organism='Artificial Sequence'.
Location/Qualifiers
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C12N15/09,C12N15/00
Description of Artificial Sequence: a

    .21
    forganism="synthetic construct"
|mol type="genomic DNA"
|db_xref="taxon:32630"

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Method for amplifying DNA.
BD183598
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/organism="synthetic construct"
/mol_type="genomic DNA"
/mol_type="genomic DNA"
/not="reacon:32630"
/note="Description of Combined DNA/RNA Molecule:Synthetic Oligomer Sequence-Synthetic Probe Sequence"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wen, X.Y., Stewart, A.K., Teui, L.C. and Hegele, R.A.
Lipase genes and proteins
Datent: Wo 0305595-A 75 10-JUL-2003;
Wen, Xiao-Yan (CA); Stewart, A., Keith (CA); Tsui, Lap-Chee (CN);
Hegele, Robert, A. (CA)
                                                                                                                                                                                                                                                                                                                        Kirtsen, N.V., Hyldig-Nielsen, J.J. and Williams, B.F.
Methods, kits and compositions pertaining to the suppression of
detectable probe binding to randomly distributed repeat sequences
in genomic nucleic acid
Patent: WO 03027328-A 19 03-APR-2003,
Boston Probes, Inc. (US); DakoCytomation Denmark A/S (DK)
Location/Qualifiers
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Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                   Sequence 19 from Patent WO03027328. AX741045
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/organism="Homo sapiens"
2280
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                   21 TTTTAGTAGAGGGGGTTTC
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artificial sequences.
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RESULT 749

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PAT 20-SEP-2000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PAT 16-MAY-2001
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1 (bases 1 to 24)

Bell,G.I., Yamagata,K., Oda,N., Kaisaki,P.J., Furuta,H.,
Horikawa,Y. and Menzel,S.
Mutations in the diabetes susceptibility genes hepatocyte nuclear factor (HNF) 1 alpha (alpha.), HNF1.beta. and HNF4.alpha
Patent: US 6187533-A 122 13-FEB-2001;
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Sequence 122 from patent US 6187533.
AR129533.
AR129533.1 GI:14117430

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                                                                                    A99115 24 bp
Sequence 19 from Patent W09909054.
A99115.1 GI:6782068
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Sequence 17 from Patent WO0115676.
AX092605
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AR129533/c
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                                                                                                                                                                                                                                                                                                                                                                                                                         Wirtz,R., Munnes,M. and Kallabis,H.
Methods and compositions for the prediction, diagnosis, prognosis,
prevention and treatment of malignant neoplasia
Patent: EP 1365034-A 244 26-NOV-2003;
Bayer HealthCare AG (DE)
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Pred. No. 8.6e+02;
0; Mismatches 1;
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/db_xref="taxon:32630"
1. .22
/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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1 (bases 1 to 20)

Hardy,J.A. and Goate,A.M.

Mutant 5182 genes

Patent: US 5973113-A 11 26-OCT-1999;

Location/Qualifiers
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Sequence 244 from Patent EP1365034.
AX938799
AX938799.1 GI:40733179
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AR082561
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/wol_type="unassigned DNA"
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                                                                                                                                                  2340 CCAAAGTGCTGGGATTACAGG 2360
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Best Local Similarity 95.2%;
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24 bp DNA linear PAT 27-AUG-2002 Peroxisome-related polypeptide, nucleotide sequence encoding the polypeptide, and utilization thereof in diagnosis and/or treatment of lung injury or disease, and diagnosis and/or treatment of oxidative stress-related disease.
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                                                                     Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                 Hayden, M.R., Brooks-Wilson, A.R., Pimstone, S.N. and Clee, S.M. Compositions and methods for modulating hdl cholesterol and triglyceride levels
Patent: WO 01156-6-A 17 08-MAR-2001;
University of British Columbia (CA); Xenon Genetics Inc. (CA)
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Cnupe, B., Erman, C., Bernard, A., Watie, R. and Farumanu, P.
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/noC="Primer"
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/organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"

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                                 Homo sapiens (human)
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A61P9/10,A61P11/06,A61P17/04,A61P19/08,A61P19/10,A61P21/00, PC
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Peroxisome-related polypeptide, nucleotide sequence encoding the polypeptide, and utilization thereof in diagnosis and/or treatment of lung injury or disease, and diagnosis and/or treatment of oxidative stress-related disease.

UNIVERSITE CATHOLIQUE DE LOUVAIN, UNIVERSITE DE MONT ZENO ON UNIQUENTIFICATHOLIQUE DE LOUVAIN, UNIVERSITE DE MONT ZENO ON UNIQUENTIFIE A 16 18-SEP-2001

PN J 2001514874-A/16

PD 18-SEP-2001

PR 20-AUG-1999 DE 9700692

PR 20-AUG-1997 DE 9700692

PR HERNARD CNUPE, CEDRICK ERMAN, ALFRED BERNARD, RUDY WATIE, PAUL PI
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CC Topology: Linear;
CC Peroxisome-riated polypeptide, nucleotide sequence encoding CC the the CC polypeptide, and utilization thereof in diagnosis and/or CC
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
mdz12
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R 20-AUG-1997 BE 9700692
I BERNARD CNUPE, CEDRICK ERMAN, ALFRED BERNARD, RUDY WATIE, PAUL PARUMANU
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/organism='Unidentified'.
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Aeomica, Inc. (US)
Location/Qualifiers
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and mdz12
                                                                                    Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Disease risk estimating method using sequence polymorphisms in a
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Best Local Similarity 87.5%; Pred. No. 9.2e+02;
Matches 21; Conservative 0; Mismatches 3;
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Aeomica, Inc. (US)
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Aeomica, Inc. (US)
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Sequence 7 from Patent WO2004003229.
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CQ760565.1 GI:44904068
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/db_xref="taxon:9606"

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    /organism="Homo sapiens"
    /mol_type="unassigned DNA"
    /db_xref="taxon:9606"

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Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                                                                      Shannon, M., Gu, Y. and Nguyen, C.T. Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Eukaryota; Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
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Aeomica, Inc. (US)
Location/Qualifiers
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                                                                                                              Sequence 5569 from Patent EP1281758.
AX692837
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AX692840
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 2086 TTATTTTTTTTGAGACCGAGT 2109
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Kieback, D.G.
Methods for diagnosing an increased risk for breast or ovarian
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tive 0; Mismatches 0; Indels
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Matches 19; Conservative 0; Mismatches 0;
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/organism="synthetic construct"
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/mol_type="unassigned"
/nofe="pximer"
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Seguence 1017 from Patent WO0129262.
AX115894
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Patent: US 5683885-A 10 04-NOV-1997;
Location/Qualifiers
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Sequence 10 from patent US 5683885.
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                                                                              2342 AAAGTGCTGGGATTACAGG 2360
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Best Local Similarity 100.0
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Disease risk estimating method using sequence polymorphisms in specific region of chromosome 19
Patent: WO 2004003229-A 133 08-JAN-2004;
Aarhus University (DK); Arbejdsmil Institute of Occupational Health) (DK)
Location/Qualifiers
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Kieback, D.G.
Methods for diagnosing an increased risk for breast or ovarian
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specific region of chromosome 19
Patent: WO 2004003229-A 7 08-JMN-2004;
Aarhus University (DK); Arbejdsmilj Instituttet (National Institute of Occupational Health) (DK)
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    .19
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    /noTe="Probe"

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Sequence 10 from patent US 5645995.
152002.1 GI:2473203
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Location/Qualifiers
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Best Local Similarity 100.
Matches 19; Conservative
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  Sequence 8 from patent US 6133503
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AR116725.1 GI:14097047
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1 (bases 1 to 20)
Scheffler, I.E.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A method of arraying genome clone
Patent: JP 2001321190-A 1518 20-NOV-2001;
THE INSTITUTE OF PHYSICAL AND CHEMICAL RESEARCH, YUGENKAISHA
GENOTECHS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Description of Artificial Sequence:Synthetic DNA FH Location/Qualifiers
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                                                                                    Picoult-Newburg, L. and Pohl, M. Genotyping reagents, kits and methods of use thereof Patent: WO 0129262-A 1473 26-APR-2001; Orchid BioSciences, Inc. (US)
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    .19
        /organism="synthetic construct"
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/db_xref="taxon:32630"

                                                                                                                                                                 /organism="synthetic construct"
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JP 2001321190-A/1518
20-NOV-2001
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AX116350
AX116350.1 GI:14033292
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JP 2001321190-A/1518.
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BD089274/c
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AR116725/c
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Unknown.
Unclassified.
I (bases 1 to 20)
Scheffler,I.B.
Mammalian artificial chromosomes and methods of using same Patent: US 613553-A 8 17-OCT-2000;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mammalian artificial chromosomes and methods of using same Patent: US 5721118-A 8 24-FEB-1998; Location/Qualifiers
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0.8%; Score 19; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 8.7e+02;
Matches 19; Conservative 0; Mismatches 0;
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Best Local Similarity 100.0%; Pred. No. 8.7e+02;
Matches 19; Conservative 0; Mismatches 0;
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1 (bases 1 to 20)
Agrawal,S. and Kandimalla,E.R.
Pseudo-cyclic oligonucleobases
Patent: US 6383752-A 23 07-MAY-2002,
Location/Qualifiers
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Sequence 23 from patent US 6383752.
AR208407. GI:21509553
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Sequence 8 from patent US 5721118.
189275
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Gaps

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PAT 26-SEP-2002

REFERENCE AUTHORS

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Unknown.
Unclassified.
1 (bases 1 to 20)
Freier,S.M.
Antisense modulation of cellular apoptosis susceptibility gene
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                 1 (bases 1 to 20)
Agrawal, S. and Kandimalla, B.R.
Pseudo-cyclic oligonucleobases
Patent: US 6383752-A 26 07-MAY-2002;
Location/Qualifiers
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Patent: US 6440737-A 17 27-AUG-2002;
Location/Qualifiers
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Sequence 1218 from Patent WO0129262.
AX116095
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Sequence 17 from patent US 6440737.
AR224472.1 GI:23333312
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AR224472/c
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 95.0%; Pred. No. 8.7e+02; iive 0; Mismatches 1; Indels
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Agrawal, S. and Kandinalla, E.R.
Pseudo-cyclic oligonucleobases
Patent: US 6383752-A 24 07-MAY-2002;
Location/Qualifiers
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Agrawal, S. and Kandimalla, B.R.
Pseudo-cyclic oligonucleobases
Patent: US 6383752-A 25 07-MAY-2002;
Location/Qualifiers
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Sequence 25 from patent US 6183752.
AR208409
AR208409.1 GI:21509556
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Sequence 24 from patent US 6383752.
AR208408
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/mol_type="unassigned DNA"
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/organism="unknown"
                                                677 GAGTGAGAACAGGTGTCACC 696
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                                                                        20 GNGTGAGAACAGGTGTCACC 1
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Best Local Similarity 95.0
Matches 19; Conservative
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AR208408/c
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TITLE
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TITLE
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
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Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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100.0%; Pred. No. 9.3e+02;
iive 0; Mismatches 0; Indels
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Polymorphisms in the human hmg-coa reductase gene
Patent: WO 0079003-A 51 28-DEC-2000;
AstraZeneca UK Limited (GB)
Location/Qualifiers
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    .25
    forganism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="taxon:32630"

                                                                                                       DNA
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AX693014
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                                                                                                    AX060516 25 bp
Sequence 51 from Patent W00079003.
AX060516

    .25
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         CACCACACCTGGCTAATTTT 2254
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Location/Qualifiers
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Homo sapiens
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Unclassified.

1 (bases 1 to 22)
Teng, D.H.-F., Tavtigian, S.V., Perry, W.L. III and Skolnick, M.H.
Specific mutations of map kinase 4 (MKK4) in human tumor cell lines identify it as a tumor suppressor in various types of cancer Patent: US 5989885-A 11 23-NOV-1999;
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                                                                              Unclassified.

1 (bases 1 to 22)

Sannay, G.A. and Smith, M.W.

Method for generation of sequence sampled maps of complex genomes

Patent: US 5851760-A 257 22-DEC-1998;
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Sequence 1197 from Patent WO0129262.
AX116074
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Sequence 11 from patent US 5989885.
AR088425.1 GI:10015188
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GI:5998131
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RESULT 785 AX817228

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DEFINITION ACCESSION VERSION KEYWORDS SOURCE

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PAT 06-AUG-2001

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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                        Ibd-related polymorphisms
Patent: WO 0142511-A 1707 14-JUN-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH (US) ; Ellipsis
Biotherapeutics Corporation (CA)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Ulfendahl, P.J. and Wong, K.C. Primers for identifying typing or classifying nucleic acids Patent: WO 0065088-A 452 02-NOV-2000; Amersham Pharmacia Biotech AB (SE)
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0.8%; Score 18.8; DB 1; Length 24;
Best Local Similarity 87.0%; Pred. No. 9.4e+02;
Matches 20; Conservative 0; Mismatches 3; Indels
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0.8%; Score 18.8; DB 1; Length 25;
Best Local Similarity 90.9%; Pred. No. 9.5e+02;
Matches 20; Conservative 0; Mismatches 2; Indels
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/organism="synthetic construct"
/mol_type="unassigned DNA"
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/note="HIA-C Homozygote Primer Sequence"
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                                                                                                                                                                                                                                                                                                                                                                                                                                              /organism="Homo sapiens"
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Sequence 452 from Patent WO0065088.
AX042886
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Sequence 12 from Patent W00172822.
AX259785.
AX259785.1 GI:16508859
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synthetic construct
artificial sequences.
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                                                                                 Gaps
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/db_xref="taxon:32630"
/note="Beschreibung der kunstlichen Sequenz: Primer i
PCR-Reaktion zur Amplifikation von M30 des Menschen"
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                                     Query Match
0.8%; Score 18.8; DB 1; Length 22;
Best Local Similarity 90.9%; Pred. No. 9.2e+02;
Matches 20; Conservative 0; Mismatches 2; Indels
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0.8%; Score 18.8; DB 1; Length 22;
Best Local Similarity 90.9%; Pred. No. 9.2e+02;
Matches 20; Conservative 0; Mismatches 2; Indels
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Patent: WO 0221138-A 40 14-MAR-2002;
Axaron Bioscience AG (DE)
Location/Qualifiers
1. .22
/organism="synthetic construct"
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IMPERIAL COLLEGE INNOVATIONS LIMITED (GB)
Location/Qualifiers
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Sequence 40 from Patent W00221138.
AX817228
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PAT 23-NOV-2000

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PAT 26-OCT-2001

FEATURES

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PAT 31-MAR-2003

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Shannon,M., Gu,Y. and Nguyen,C.T.
Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
mdz12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                  Query Match 0.8%; Score 18.8; DB 1; Length 25; Best Local Similarity 90.9%; Pred. No. 9.5e+02; Matches 20; Conservative 0; Mismatches 2; Indels
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CARBOHYDRATE-DEFICIENT GLYCOPROTEIN SYNDROME TYPE I
PATENT: WO 9849324-A 13 05-NOV-1998;

MATTHIJS GERT (BE); GENZYME LTD (GB)

Location/Qualifiers
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Aeomica, Inc. (US)
Location/Qualifiers
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Seguence 5720 from Patent EP1281758.
AX692988
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Sequence 13 from Patent WO9849324.
A83584
A83584.1 GI:6732840
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1 (bases 1 to 20)
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Best Local Similarity 90.9
Matches 20; Conservative
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                                                                                           Hugot,J.P., Thomas,G., Zouali,M., Lesage,S. and Chamaillard,M.
Genes involved in intestinal inflammatory diseases and use thereof
Patent: W0 0172822-A 12 04-OCT-2001;
Fondation Jean Dausset-Ceph (FR)
Location/Qualifiers
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and mdz12
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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                                  Bukaryoča, Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Aeomica, Inc. (US)
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AX692915
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Sequence 5568 from Patent EP1281758.
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Homo sapiens (human)
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RESULT 794 A83598 DEFINITION ACCESSION VERSION KEYWORDS SOURCE

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Unknown.
Unclassified.

1 (bases 1 to 20)

1 object.

1 bong.J.-T., Barrett,J.Carl., Lamb,P.W. and Isaacs,J.T.

Dong.J.-T., barrett,J.Carl., Lamb,P.W. and Isaacs,J.T.

Diagnostic methods and gene therapy using reagents derived from the human metastasis suppressor gene KAII

Patent: US 6204000-A 12 20-MAR-2001;

Location/Qualifiers
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Miragila, L. J., Nero, P., Graham, M.J. and Monia, B.P.
Antisense oligonucleotide modulation of human mdm2 expression
Patent: US 6238921-A 27 29-MAY-2001;
Location/Qualifiers
                                                                                                                                           Query Match 0.8%; Score 18.4; DB 1; Length 20; Best Local Similarity 95.0%; Pred. No. 9.2e+02; Matches 19; Conservative 0; Mismatches 1; Indels
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       Kern,S.E. and Hahn,S.A.
Tumor suppressor gene, DPC4
Patent: US 595x392-A 70 21-SEP-1999;
Location/Qualifiers
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AR142729
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Sequence 27 from patent US 6238921.
AR154610 GI:15122663
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E 1 (bases 1 to 20)

RS Kern,S.E. and Hahn,S.A.

DPC4 polypeptide

NAL Patent: US 5814457-A 70 29-SEP-1998;

Location/Qualifiers

Location/Qualifiers
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Sequence 70 from patent US 5814457.
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Sequence 27 from Patent WO9849324.
A83598.1 GI:6732854
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DEFINITION ACCESSION VERSION KEYWORDS SOURCE

RESULT 796

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RESULT 799 CQ755665/c LOCUS DEFINITION ACCESSION ö

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Sequence: an artificially
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                    Ota,T., Nishikawa,T., Isogai,T., Hayashi,K., Ishii,S., Kawai,Y.,
Wakamatsu,A., Sugiyama,T., Nagai,K., Kojima,S., Otsuki,T. and
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Polymorphisms in th clcn7 gene as genetic markers for bone mass Patent: WO 2004046381-A, 7 03-JUN-2004;
The University Court of The University of Aberdeen (GB)
Location/Qualifiers
                                                                                            Primers for synthesizing full length cDNA clones and their use
Patent: BP 1396543-A 4221 10-MAR-2004;
Research Association for Biotechnology (JP)
Location/Qualifiers
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/note="Description of Artificial
synthesized primer se q uence"
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llarity 95.0%; Pred. No. 9.2e+02;
Conservative 0; Mismatches 1;
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Patent: WO 2004033723-A 128 22-APR-2004;
IMPERIAL COLLEGE INNOVATIONS LIMITED (GB)
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Seguence 7 from Patent WO2004046381.
CQ819694

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A method for the stimultaneaous production of multiple proteins;
vectors and cells for use therein
Patent: WO 2003106684-A 172 24-DEC-2003;
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Means and methods for regulating gene expression
Patent: WO 2003106674-A 166 24-DEC-2003;
Chromagenics B.V. (NL)
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0.8%; Score 18.4; DB 1;
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/organism="synthetic construct"
/mol_type="unassigned DNA"
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/noTe="oligonucleotide E23"
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                                              Sequence 166 from Patent WO2003106674. CQ755665.1 GI:44846470
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Sequence 4221 from Patent EP1396543.
CQ784081. GI:45538569
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Sequence 70 from patent US 5712097.
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Length polymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same
Patent: US 5582979-A 341 10-DEC-1996;
Location/Qualifiers
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Leonard, W.J., Noguchi, M. and McBride, O. Wesley.
Methods for diagnosis of XSCID and kits thereof
Patent: US 5518880-A 25 21-MAY-1996;
'organism="synthetic construct"
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Sequence 341 from patent US 5582979.
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Sequence 25 from patent US 5518880.
121054
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I (bases 1 to 20)
Weston, B.W. and Hiller K.M.
Antisense human fucosyltransferase sequences and methods of use
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1 (bases 1 to 20)

Monia_B.P., Gasarde,W.A., Freier,S.M. and Wancewicz,E.
Antiesnse modulation of TERT expression
Patent: US 6492171-A 82 10-DEC-2002;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                         Query Match 0.8%; Score 18.4; DB 1; Length 20; Best Local Similarity 95.0%; Pred. No. 9.2e+02; Matches 19; Conservative 0; Mismatches 1; Indels
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1 (bases 1 to 20)

Kern,S.E. and Hahn,S.A.

Tumor suppressor gene, DPC4

Patent: US 5712097-A 70 27-JAN-1998;

Location/Qualifiers
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Location/Qualifiers
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Sequence 82 from patent US 6492171.
AR266075.1 GI:29694921
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Sequence 18 from patent US 6350868.
AR195440
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/wol_type="genomic DNA"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Polymorphic kinase anchor proteins and nucleic acids encoding the
                                                                                                                                                                                                                                                                       Braun, A., Koester, H., van den Boom, D., Ping, Y., Rodi, C., He, L.,
                                                                                                                                                                                                                                                                               Chiu, N. and Jurinke, C. Methods for generating databases and databases for identifying polymorphic genetic markers Patent: WO 012785-A 53 19-APR-2001;
                               Gaps
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    Length 20;
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Best Local Similarity 95.0%; Pred. No. 9.2e+02;
Matches 19; Conservative 0; Mismatches 1;
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Pred. No. 9.2e+02;
0; Mismatches 1;
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/note="Oligonucleotide Primer"
                                                                                                                                                                                                                                                                                                                                                                            Patent: WO 0204489-A 9 17-JAN-2002;
                                                                                                                                           Sequence 53 from Patent WO0127857.
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                                                                                                                                                                                                                                                                                                                                     Sequenom, Inc. (US)
Location/Qualifiers
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Location/Qualifiers
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Best Local Similarity
Matches 19; Conserv
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BD128005 20 bp DNA linear PAT 18-SEP-2002 Primer for synthesizing full-length cDNA and use thereof. BD128005.1 GI:23222950 JP 2002017375-A/3436.
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  PAT 25-JUN-2003
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Methods and compositions for the prediction, diagnosis, prognosis,
prevention and treatment of malignant neoplasia
Patent: EP 1365034-A 259 26-NOV-2003;
Bayer HealthCare AG (DE)
Location/Qualifiers
                                                                                                                                                                                                  Otte,A.P. and Kruckeberg,A.L.

Dna sequences comprising gene transcription regulatory qualities and methods for detecting and using such dna sequences

Patent: WO 03004704-A 169 16-JAN-2003;

Chromagenics B.V. (NL)

Location/Qualifiers
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/db xref="texon:32630"
/noTe="oligonucleotide E23"
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20 bp Dl
Seguence 169 from Patent WO03004704.
AX764699
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Sequence 259 from Patent EP1365034.
AX938814
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Libaes 1 (bases 1 to 20)
ATTHORS Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.
TITLE Antisense modulation of human MDM2 expression
TOURNAL Patent:JP 2002508944-A 27 26-MAR-2002;
SSY DIAGENTIFIED

NO Unidentified
PN JP 2002508944-A/27
PD 26-MAR-2002
PF 26-MAR-1999 JP 27
PR 26-MAR-1999 JP 27
PR 26-MAR-175
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LOREN J MIRAGLIA, PAMELA NERO, MARK J GRAHAM, BRETT P MONIA, LEX M
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                                                                                                  Primer for synthesizing full-length cDNA and use thereof Patent: JP 2002017375-A 3436 22-JAN-2002;
HELIX RESEARCH INSTITUTE
OS Unidentified
PN JP 2002017375-A/3436
PD 22-JAN-2002
PF 07-JUL-2000 JP 200253172
PT TOSHIO OTA, TETSUO NISHIKAWA, TAKAO ISOGAI, KOJI HAYASHI, SHIZUKO
                                                                                                                                                                                                                                                                                                                                         PC C12P21/02,C1201/68//C12P21/08,G06F17/30,C12N15/00,C12N5/00 CC Description of Artificial Sequence: an artificially CC synthesized primer
                                                                                                                                                                                                                                                                                            PC
C12N15/09, C07K14/47, C07K16/18, C12N1/15, C12N1/19, C12N1/21, C12N5/ PC
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C12N15/00
Strandedness: Single;
                                       1 (bases 1 to 20)
Ota,T., Nishikawa,T., Isogai,T., Hayashi,K., Ishii,S., Kawai,Y.,
Wakamatsu,A., Sugiyama,T., Nagai,K., Kojima,S., Otsuki,T. and
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PI YURI KAWAI,AI WAKAWATSU,TOMOYASU SUGIYAWA,KEIICHI NAGAI,
SHINICHI KOJIMA,
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    .20
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BD138101/c
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Aethod for distinguishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein.
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B31629 and primer to be used therein.
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/organism='Unidentified'.
Location/Qualifiers
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/organism='Artificial
Location/Qualifiers
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Pred. No. 9.4e+02;
0; Mismatches 1;
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                                                                                                                                                                  1695 TTTACATGTGCAAAGAAGCT 1714
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                                                     2351 GGATTACAGGCATGAGCCAC 2370
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synthetic construct
artificial sequences.
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     Matches 19; Conservative
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TITLE
JOURNAL
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S 2100023671-A/3.

S 37theric construct

Syntheric construct

Syntheric construct

attificial sequences.

To (bases 1 to 21)

Stohiro, O., Individual based on PCR finger print with the use of restriction primer of inter-SINE sequences

And primer to be used therein

Patent: JP 2000023671-A/3

Attificial Sequence

NATIONAL RESEARCH INSTITUTE OF AQUACULTURE

OS Artificial Sequence

PR 10-JUL-1998 JP 1998195692

PR 10-JUL-1998 JP 1998195692
                 artificial sequences.

It (bases 1 to 21)

Ichiro,O., Ichiro,N. and Hiroshi,Y.

Method for distinguishing eucaryctic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein.

A print with the use of restriction primer of inter-SINE sequences and primer to be used therein.

A print with the use of restriction primer of inter-SINE sequences and primer to be used therein.

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A print with the use of restriction primer of inter-SINE sequence.
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/organism='Artificial Sequence'.
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/organism='Artificial Sequence'.
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/db_xref="taxon:32630"

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    forganism="synthetic construct"
|mol_type="genomic DNA"
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  synthetic construct
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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/note="'n' represents a polymorphic base"
  linear
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                                                                                                                                                                                                                                                                                                                            1. .21
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21 bp Dl
Seguence 1202 from Patent WO0129262.
AXI16079
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Seguence 315 from Patent WO0134840.
AX146124
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22 bp DNA linear PAT 18-JUN-2001
Method for distinguishing evoaryotic individual based on PCR finger
print with the use of restriction primer of inter-SINE sequences
and primer to be used therein.
E31632.
E31632.1 GI:13018542
JP 2000023671-A/5.
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JP 2000023671-A/4.
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Best Local Similarity 95.0
Matches 19; Conservative
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Synthetic construct

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Method for distinguishing eucaryotic individual based on PCR finger
print with the use of restriction primer of inter-SINE sequences
and primer to be used therein.
E31634.1 G1:13018544
B21634.1 G1:13018544
B2 2000023671-A/7.
Synthetic construct
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E31633 1 GI:13018543

JP 2000023671-A/6.
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/organism='Artificial Sequence'
Location/Qualifiers
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1. 22
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Is Ichiro, 0. Ichiro, N. and Hiroshi, Y.

E 1 (bases 1 to 22)

Is Ichiro, 0. Ichiro, N. and Hiroshi, Y.

Method for distinguishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINB sequences and primer to be used therein

Artificial Sequence

NATIONAL RESEARCH INSTITUTE OF AQUACULTURE

OS Artificial Sequence

PN JP 2000023671-A/8

PP 10-JUL-1998 JP 1998195692

PR 10-JUL-1998 JP 1998195692
Method for distinguishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein primer of inter-SINE sequences and primer to be used therein primer of inter-SINE sequences and primer to be used therein primer of inter-SINE sequences.

M. Patent: JP 2000033671-A 7 25-JAN-2000;

NATIONAL RESEARCH INSTITUTE OF AQUACULTURE
OS Artificial Sequence
PN JP 2000023671-A/7

PN JP 2000023671-A/7

PN JP 2000023671-A/7

PR 10-JUL-1998 JP 1998195692

PR 10-JUL-1998 JP 1998195692

PR CIZNIS/09, CIZOII/68, CIZNIS/00

CC CIZNIS/09, CIZOII/68, CIZNIS/00

CC HK Key Location/Qualifiers

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95.0%; Pred. No. 9.5e+02;
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/organism="synthetic construct"
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JP 2000023671-A/8.
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B31637.1 GI:13018547

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S 1 (bases 1 to 2)

Method for distinguishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein.

D PATENT. JP 2000023671-A 10 25-JAN-2000;

NATIONAL RESEARCH INSTITUTE OF AQUACULTURE

OS Artificial Sequence

PN JP 2000023671-A/10

PP 10-JUL-1998 JP 1998195692

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Synthetic construct
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1 (bases 1 to 22)
Ichiro, O. Ichiro, and Hiroshi, Y.
Ichiro, C. Ichiro, Shang eucaryotic individual based on PCR finger
print with the use of restriction primer of inter-SINE sequences
and primer to be used therein
Patent: JP 2000023671-A 9 25-JAN-2000;
NATIONAL RESEARCH INSTITUTE OF AQUACULTURE
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organism='Artificial Sequence'.
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C12N15/09, C12Q1/68, C12N15/00
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Best Local Similarity 95.0%; Pred. No. 9.5e+02;
Matches 19; Conservative 0; Mismatches 1;
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    .22
    forganism="synthetic construct"
|mol type="genomic DNA"
|db_xref="taxon:32630"

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10-JUL-1998 JP 1998195692
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JP 2000023671-A/9
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   E31639.1 GI:13018549 JP 2000023671-A/12.
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AR154030/c
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DEFINITION 8
 VERSION
KEYWORDS
SOURCE
ORGANISM
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ACCESSION
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AX609024/c
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/organism='Artificial Sequence'.
Location/Qualifiers
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                                              ce 1. .22 /organism='Artificial Sequence'
Location/Qualifiers
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ICHIRO OHARA,ICHIRO NAKAYAMA,HIROSHI YASUE
C12N15/09,C12Q1/68,C12N15/00

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    ^coganism="synthetic construct"
|mol_type="genomic DNA"
|db_xref="taxon:32630"

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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Best Local Similarity 95.03
Matches 19; Conservative
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artificial sequences.

I (bases 1 to 22)

I (bases 2 to 22)

I (bases 3 to 22)

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Catarrhini, Hominidae, Homo.
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Location/Qualifiers
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/organism="synthetic construct"
/wol_type="genomic DNA"
/db_xref="reaxon:32630"
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Coronary chip
Patent: WO 02072882-A 49 19-SEP-2002;
OGHAM GmbH (DE)
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Sequence 49 from Patent WO02072882.
AX609024
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Mammalia; Eutheria; Primates;
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Query Match
Best Local Similarity 89.5'
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Unclassified.
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I Patent: JP 2002502606-A 80 29-JAN-2002;
PROMEGA CORP
OS Unidentified
PN JP 2002502666-A/80
PD 29-JAN-2002
PP 04-FEB-1999 UP 200530608
PR 04-FEB-1999 UP 209/018584
PI JAMES W SCHUMM, JEPFREY W BACHER
PC CLINIS/09, Cli201/68, Clinis/09, Cli201/68, Linear;
CC Strandedness: Single;
CC Material and method for specifying and analyzing medium-size
CC CNA material and method for specifying and analyzing medium-size
                                                                                                                                                                                                                                                                                                                                    Material and method for specifying and analyzing medium-size tandem BD130136
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JP 2002502606-A/80.
unidentified
unclassified.
1 (Dases I to 24)
Schumm, J.W. and Bacher, J.W.
Material and method for specifying and analyzing medium-size tandem
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                                                             Unclassified.

1 (bases 1 to 24)
Schumm, J.W. and Bacher, J.W.
Schumm, J.W. and methods for indentifying and analyzing intermediate tandem repeat DNA markers
Patent: US 6238863-A 80 29-MAY-2001;
Location/Qualifiers
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/organism='Unidentified'
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Location/Qualifiers
                                                                                                                                                         1. .24
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    .24
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              GI:15122083
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 AR154030
AR154030.1
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RESULT 832

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PAT 28-AUG-2000
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DNA sequence encoding the myotonic dystrophy gene and uses thereof
Patent: US 5977333-A 13 02-NOV-1999;
Location/Qualifiers
                                                                                                                                                          1 (bases 1 to 19)
Brook,J.David., Housman,D.E., Shaw,D.J., Harley,H.G. and Johnson,K.J.
DNA sequence encoding the myotonic dystrophy gene and uses thereof Patent: US 5955265-A 13 21-SEP-1999;
Location/Qualifiers
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Unknown.
Unclassified.
1 (bases 1 to 19)
Siciliano,M.J. and Liu,P.
Siciliano,M.J. and Liu,P.
In-situ hybridization probes for identification and banding of specific human chromosomes and regions
Patent: US 5538669-A 1 23-ULL-1996;
Location/Qualifiers
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Brook,J.David., Housman,D.E., Shaw,D.J., Harley,H.G. and
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89.5%; Pred. No. 9.3e+02;
tive 2; Mismatches 0; Indels
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Sequence 13 from patent US 5977333.
AR083935.1 GI:10010706
19 bp
Sequence 13 from patent US 5955265.
AR074596
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/mol_type="unassigned DNA"
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Sequence 1 from patent US 5538869.
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123815.1 GI:1603685
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FEATURES
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PAT 29-MAR-1999

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Gaps

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PAT 03-MAR-2004

linear

RESULT 835

AUTHORS REFERENCE

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Unknown.
Unclassified.
1 (bases 1 to 23)
MacKenzie,A.E., Korneluk,R.G., Roy,N., Mahadevan,M.S., McLean,M.
                                                                                                                                                                  Korneluk, R.G., Mackenzie, A.E., Roy, N., Robertson, G. and Tamai, K. USE OF NBURONAL APOPTOSIS INHIBITOR PROTEIN (NAIP)
PAtent: WO 9726331-A 16 24-UUL-1997;
UNIV OTTAWA. (CA)
Other publication AU 1614997 19970811.
                                                                                                                                                                                                                                                                                                                                                        Query Match 0.8%; Score 18.2; DB 1; Length 23; Best Local Similarity 87.0%; Pred. No. 9.8e+02; Matches 20; Conservative 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Patent: WO 2004005547-A 137 15-JAN-2004, IMPERIAL COLLEGE INNOVATIONS LIMITED (GB) Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.23
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Seguence 137 from Patent WO2004005547.
CQ766176
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Sequence 16 from patent US 6429011.
AR222091.1 GI:23329461
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Sequence 16 from Patent WO9726331.
A64524
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synthetic construct
artificial sequences.
                                                                A64524.1 GI:3717923
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unidentified
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Matches 20; Conserv
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AR222091/c
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CQ766176
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A novel nucleic acid molecule
Patent: WO 9851790-A il9-NOV-11998;
CANCILLA MICHAEL ROBERT (AU) ; CHOO KONG HONG ANDY (AU) ; SART
DESIREE DU (AU) ; AMRAD OPERATIONS PTY LTD (AU)
Location/Qualifiers
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0.8%; Score 18.2; DB 1; Length 19;
Best Local Similarity 89.5%; Pred. No. 9.3e+02;
Matches 17; Conservative 2; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                Unclassified.

1 (bases 1 to 19)
Gilliam, T.Conrad. and Tanzi, R.E.
Wilson's disease gene
Patent: US 5578493-A 1 26-NOV-1996;
Location/Qualifiers
1. .19
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/mol_type="unassigned DNA"

    .19
/organism="unidentified"
/mol_type="unassigned DNA"
/db_xref="taxon:32644"

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Sequence 1 from patent US 5578493.
129969
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Sequence 1 from Patent W09851790.
AX033909
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RESULT 836

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AUTHORS REFERENCE

RESULT 837

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Gaps

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Indels

Length 23;

PAT 26-SEP-2002

linear

PAT 14-JAN-2004

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BD130152.1 GI:23225097
JP 20025052606-A/96.
unidentified
unclassified.
unclassified.
1 (bases I to 24)
Schumm, J.W. and Bacher, J.W.
Material and method for specifying and analyzing medium-size tandem
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Material and method for specifying and analyzing medium-size tandem
ED130152
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 24;
                                                                                                                                                                                                                                                                                          Immunostimulatory oligonucleotides and uses thereof Patent: WO 03101375-A 33 11-DEC-2003; IMMUNOTECH S.A. (AR) Location/Qualifiers
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Pred. No. 9.9e+02;
0; Mismatches 3; Indels
                                                                                                                                  linear
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Location/Qualifiers
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PRECENT: UP 2002502606-A 96 29-JAN-2002;
PROMEGA CORP
OS Unidentified
PN JP 2002502606-A/96
PP 29-JAN-2002
PF 04-FEB-1999 JP 2000530608
PR 04-FEB-1999 US 09/018584
PR JAMES W SCHUMM, JEFFREY W BACHER
PC CL2M15/09, C12Q1/68, C12M15/00
CC Strandedness: Single;
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Sequence 33 from Patent W003101375.
AX961638
AX961638.1 GI:40881096
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                    2322 TCCGCCCACCTCGGCCTCCCAAAG 2345
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1 (bases 1 to 24)
Schumm,J.W. and Bacher,J.W.
Materials and methods for indentifying and analyzing intermediate tandem repeat DNA markers
Patent: US 6238863-A 96 29-MAY-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
             Neuronal apoptosis inhibitor protein gene sequence and mutations causative of spinal muscular atrophy Patent: US 6429011-A 16 06-AUG-2002; Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Daly,M., Hudson,T.J., Lander,E.S., Rioux,J. and Siminovitch,K. Ibd-related polymorphisms
Patent: WO 042511-A 1887 14-JUN-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH (US); Ellipsis Biotherapeutics Corporation (CA)
Location/Qualifiers
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                                                                                                                                          0.8%; Score 18.2; DB 1; Length 23;
87.0%; Pred. No. 9.8e+02;
7ative 0; Mismatches 3; Indels
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Query Match

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Matches 20; Conservative 0; Mismatches 3;
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Sequence 96 from patent US 6238863.
ARIS4046.1 GI:15122099

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                                                                                             /organism="unknown"
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Thes 20; Conservative
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and Ikeda, J.-E.
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AX184134/c
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PAT 18-DEC-2003

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BD09344.1 GI:2639032

BD093444.1 GI:2639032

JP 2001527419-A/6.

Synthetic construct

Synthetic construct

artificial sequences.

The (bases 1 to 18)

Almemii.E.S.

FADD-like anti-apoptotic molecules, methods of using the same, and compositions for and methods of making the same

AL Patent: JP 2001527419-A 6 25-DEC-2001;

THOMAS. USFPERSON UNIVERSITY

PN JP 2001527419-A/6

PD 25-DEC-2001

PP 20-MAY-1998 JP 1998550515
                                                                                                                       PADD-like anti-apoptotic molecules, methods of using the same, and compositions for and methods of making the same Patent: US 6576751-A 9 10-UNN-2003; Location/Qualifiers
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BD093444
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1 (bases 1 to 18)
DePinho,R.A.
Method of inhibiting cell proliferation using an anti-oncogene
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0.8%; Score 18; DB 1; Length 18;
Best Local Similarity 100.0%; Pred. No. 9.3e+02;
Matches 18; Conservative 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Pred. No. 9.3e+02;
Matches 18; Conservative 0; Mismatches 0;
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Patent: US 6613750-A 2 02-SEP-2003;
Location/Qualifiers
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Sequence 2 from patent US 6613750.
AR392161
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/mol_type="genomic DNA"
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 AR343036.1 GI:33738354
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Alnemri, E.S.
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Alnemri,E.S.
FADD-like anti-apoptotic molecules, methods of using the same, and compositions for and methods of making the same
Patent: US 6207801-A 9 27-MAR-2001;
Location/Qualifiers
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Unclassified.
1 (bases 1 to 18)
Monia, B.P., Baker, B.F. and Cowsert, L.M.
Antisense modulation of CREL expression
Patent: US 6001652-A 30 14-DEC-1999;
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Sequence 30 from patent US 6001652.
AR094528
AR094528.1 GI:10021535
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AR140525
AR140525.1 GI:14483021
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Sequence 9 from patent US 6576751.
AR343036
/db_xref="taxon:32644"
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Matches 18; Conservative
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DEFINITION

RESULT 845

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VERSION KEYWORDS SOURCE ORGANISM

REFERENCE AUTHORS TITLE

JOURNAL FEATURES

AR343036 LOCUS DEFINITION ACCESSION

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PAT 15-MAY-2001

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Secreted human proteins, polynucleotides encoding them and methods of using the same Patent: WO 0119856-A 37 22-MAR-2001; Curagen Corporation (US)
                                                                                                                                                                                                                                                                                                Shimkets, R.A., Fernandes, E., Herrmann, J.L., Liu, X., Yang, M. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Korneluk, R.G., Lacasse, E., Baird, S., Holcik, M. and Young, S. Antisense iap nucleic acids and uses thereof
Patent: WO 0226958-A 98 04-APR-2002;
University of Ottawa (CA); Aegera Therapeutics Inc. (CA)
Location/Qualifiers
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100.0%; Pred. No. 9.4e+02;
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          0; Mismatches
                                                                                                                                                    AX133851 19 bp
Sequence 37 from Patent WO0119856.
AX133851
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Sequence 98 from Patent WO0226968.
AX411998
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                                        2124 CAGGCTGGAGTGCAGTGG 2141
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          Matches 18; Conservative
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Best Local S:
Matches 18
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BD143839
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20-MAY-1997 US 08/859167
EMAD S ALNEMRI
CO7H11/04,G01N33/48,G01N33/53,G01N33/574,C12P21/06,C07K16/00
Strandedness: Single;
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Korneluk, R.G., LaCasse, E., Baird, S., Holcik, M. and Young, S. Antisense IAP nucleic acids and uses thereof

Patent: US 6673917-A 98 06-JAN-2004;

Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.8%; Score 18; DB 1; Length 19; Best Local Similarity 100.0%; Pred. No. 9.4e+02; Matches 18; Conservative 0; Mismatches 0; Indels
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Best Local Similarity 100.0%; Pred. No. 9.3e+02;
Matches 18; Conservative 0; Mismatches 0;
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db_xref="teaxon:32630"
/noTe="Primer"

    18
    forganism="synthetic construct"
/mol type="genomic DNA"
    /db_xref="taxon:32630"

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Sequence 98 from patent US 6673917.
AR451453
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Sequence 106 from Patent WO0129262.
AXII4983
AXII4983.1 GI:14031925
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/organism="unknown"
/mol_type="genomic DNA"
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                                                                  Topology: Linear;
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PAT 14-JUN-2002

ö Gaps ö query match 0.8%; Score 18; DB 1; Length 19; Best Local Similarity 100.0%; Pred. No. 9.4e+02; Matches 18; Conservative 0; Mismatches 0; Indels

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Synthetic construct DNA, forward primer for human STS sts-A009X34
at 1p36.
     PAT 17-JAN-2003
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Watanabe,N., Inazawa,J., Hosoda,F., Arai,Y., Mizushima,H.,
Morohashi,A., Ohira,M., Nakagawara,A., Liu,S., Hoshi,M., Horii,A.
                                                                                synthetic construct
artificial sequences.

1 (bases 1 to 19)
Sugita,Y., Hashida,R., Ogawa,K., Obayashi,M., Nagasu,T. and
Tsujimoto,K.
Method of examining allergic disease
Patent: JP 2002095500-A 7 02-APR-2002;
GENOX PRESEARCH INC,THE DIRECTOR OF NATIONAL CHILDREN'S HOSPITAL
OS Artificial Sequence
PN JP 2002095500-A/7
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PC C12Q1/02,G01N33/15,G01N33/50//C12P21/08,C12N5/00,C12N5/00,
                                                                                                                                                                                                                                                                   02-APR-2002
25-SEP-2000 JP 200291316
YUJI SUGITA,RYOICHI HASHIDA,KAORU OGAWA,MASAYA OBAYASHI,
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BAC-based STS-content map spanning a 35-Mb region of human
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Location/Qualifiers
      linear
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100.0%; Pred. No. 9.4e+02;
tive 0; Mismatches 0; Indels
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
Method of examining allergic disease.
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Genomics 74 (1), 55-70 (2001)
21269192
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                                                   BD143839.1 GI:27849597 JP 2002095500-A/7.
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2 (bases 1 to 19)
Horii, A.
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                                                                                                                                  /organism="synthetic construct"
/mol_type="genomic DNA"
/mol_type="taxon:32630"
1. 19
/note="forward primer for human STS sts-A009X34 at 1p36
sts-A009X34 obtained from clones B61B17, B86A23, B26B112,
B316H11, B26P17, Human BAC library RPCI-11"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Platzer, M., Platzer, C., Gudermann, T., Hebebrand, J., Hinney, A. and Reichwald, K.
Mchri variant associated with human obesity
Patent: WO 200104489-A 57 18-DEC-2003;
Philipps-Universitaet Marburg (DE)
Location/Qualifiers
Direct Submission
Submitted (04-AUG-2001) Akira Horii, Tohoku University School of
Medicine, Molecular Pathology; 2-1 Seiryomachi, Aoba-ku, Sendai,
Miyagi 980-8575, Japan (E-mail:horii@mail.cc.tohoku.ac.jp,
Tel:81-22-717-8047)
Location/Qualifiers
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1 (bases 1 to 20)

McKay.R., Freier.S.M. and Wyatt,J.

Antisense modulation of casein kinase 2-alpha prime expression patent: US 6455307-A 18 24-SEP-2002;

Location/Qualifiers
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100.0%; Pred. No. 9.4e+02;
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100.0%; Pred. No. 9.6e+02;
ative 0; Mismatches 0;
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/nofe="pximer ABr"
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Sequence 18 from patent US 6455307.
AR23228.1 GI:27274220
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Matches 18; Conservative
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Unthingue.
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Unclass 1 to 21)
Munroe, D.J. and Housman, D.E.
Interspersed repetitive element-bubble amplification of nucleic acids
acids
Patent: US 5597694-A 2 28-JAN-1997;
Location/Qualifiers
                                                                                      Sasagawa,T.
Method for identifying HPV infection type
Patent: JP 200133168-A 68 20-NOV-2001;
TOSHIYUKI SASAGAWA
OS Artificial Sequence
PN JP 2001331168-A/68
PN JP 2001321168-A/68
PN JP 20010321168-A/68
PN JP 20004-2001
PF 12-NAX-2000 JP 2000140602
PI TOSHIYUKI SASAGAWA
CC TINIS/09,C1201/68//G01N33/569
CC TINIS/09,C1201/68//G01N33/569
CC TINIS/O9 C1201/68//G01N33/569
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/db_xrefe"taxon:32630"
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Sequence 87 from patent US 6475739.
AR242941
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Sequence 2 from patent US 5597694.
I34288.1 GI:1825079
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 BD085495.1 GI:22631105
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synthetic construct
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Unknowm.
Unclassified.
1 (bases 1 to 22)
An,G., O'Hara,S.Mark., Ralph,D. and Veltri,R.
Biomarkers and targets for diagnosis, prognosis and management of
prostate, breast and bladder cancer
Patent: US 6218529-A 87 17-APR-2001;
Location/Qualifiers
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( Dages 1 to 20)

Watt,A.T.

Antisense modulation of caspase 6 expression

Patent: US 6566135-A 4 20-MAY-2003;

Location/Qualifiers
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Best Local Similarity 100.0%; Pred. No. 9.9e+02;
Matches 18; Conservative 0; Mismatches 0;
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Sequence 87 from patent US 6218529.
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Sequence 4 from patent US 6566135.
AR337079
AR337079.1 GI:33722933
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/mol_type="genomic DNA"
/mol_type="genomic DNA"
                                                                                         2124 CAGGCTGGAGTGCAGTGG 2141
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Gaps

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Key

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DNA

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Gaps

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PAT 20-DEC-2002

linear

DNA

/note="Description of Combined DNA/RNA Molecule:Synthetic

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Query Match
Best Local Similarity
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Methods, kits and compositions pertaining to the suppression of
detectable probe binding to randomly distributed repeat sequences
in genomic nucleic acid
Patent: WO 03027328-A 6 03-APR-2003,
Boston Probes, Inc. (US); DakoCytomation Denmark A/S (DK)
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Methods for identifying genomic deletions
Patent: WO 0210455-A 87 07-FEB-2002;
Celltech R & D, Inc. (US) ; Straehling-Hampton, Karen (US)
                              1 (bases 1 to 21)
Brunkow, M.E., Proll, S., Paeper, B. and Staehling-Hampton, K.
Methods for identifying genomic deletions
Patent: US 6475739-A 87 05-NOV-2002,
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Sequence 6 from Patent W003027328.
AX741032.
AX741032.1 GI:30523893
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AX384993
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/note="Description of Combined DNA/RNA Molecule:Synthetic Oligomer Sequence-Synthetic Probe Sequence"
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Methods, kits and compositions pertaining to the suppression of
detectable probe binding to randomly distributed repeat sequences
in genomic nucleic acid
Patent: WO 03027328-A 18 03-APR-2003;
Boston Probes, Inc. (US); DakoCytomation Denmark A/S (DK)
Location/Qualifiers
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methods, kits and compositions pertaining to the suppression of
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detectable probe binding to randomly distributed repeat sequences
in genomic nucleic acid
Patent: Wo 03027328-A 25 03-APR-2003;
Boston Probes, Inc. (83); pakoCytomation Denmark A/S (DK)
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0.8%; Score 17.8; DB 1; Length 21;
Best Local Similarity 90.5%; Pred. No. 9.9e+02;
Matches 19; Conservative 0; Mismatches 2; Indels
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                                            0.8%; Score 17.8; DB 1; Length 21; larity 90.5%; Pred. No. 9.9e+02; Conservative 0; Mismatches 2; Indels
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AX741051
AX741051.1 GI:30523912
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Sequence 18 from Patent W003027328.
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                                                                               Best Local Similarity
Matches 19; Conserv
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Matches

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KEYWORDS SOURCE

REFERENCE AUTHORS TITLE

JOURNAL

PEATURES

ACCESSION VERSION

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PAT 11-MAY-2001
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Test system for the discovery of active agents in nerve cell
diseases
                                                                                                                                                                       Patent: WO 2004020661-A 38 11-MAR-2004;
MedInnova Gesellschaft fuer Medizinische Innovation en aus
Akademischer Forschung mbH (DE)
Locar Forschung fiers
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1 (bases 1 to 19)
Brook.J.David., Housman,D.E., Shaw,D.J., Harley,H.G. and Johnson,K.J.
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Patent: WO 0129262-A 2074 26-APR-2001;
Orchid BioSciences, Inc. (US)
Location/Qualifiers
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/db xref="taxon:32630"
/noTe="Primer"
                                                                                                                                                                                                                                                           /organism="synthetic construct"
/mol_type="unassigned DNA"
/mol_type="taxon:32630"
/nofe="primer"
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Sequence 2074 from Patent WO0129262.
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Sequence 14 from patent US 5955265.
AR074597.1 GI:10001350
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 CQ787032.1 GI:45722015
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Bukaryota, Viridiplantae, Streptophyta, Embryophyta, Tracheophyta,
Spermatophyta, Coniferopsida, Coniferales, Pinaceae, Abies.
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 2; Indels
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Cd2000 and cd2001 molecules and uses thereof
Patent: EP 1223218-A 23 17-JUL-2002;
Millennium Pharmaceuticals, Inc. (US)
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                 1. .21
/organism="synthetic construct"
/mol_type="unassigned DNA"
/db xref="taxon:32630"
/noTe="Primer"
                                                                                                                                                       DNA
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Patent: WO 03050301-A 89 19-JUN-2003;
Gurling, Hugh Malcolm Douglas (GB)
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 0; Mismatches
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Sequence 89 from Patent WO03050301.
AX786478.1 GI:32953098

    22
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Sequence 23 from Patent BP1223218.
AX474262
                              2117 TGTTACCCAGGCTGGAGTGCA 2137
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CQ787032/c
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AX474262/c LOCUS DEFINITION

RESULT 867

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DNA sequence encoding the myotonic dystrophy gene and uses thereof Patent: US 5955265-A 14 21-SEP-1999, Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 19)
Brook, J. David., Housman, D.E., Shaw, D.J., Harley, H.G. and Johnson, K.J.
DNA sequence encoding the myotonic dystrophy gene and uses thereof Patent: US 5977333-A 14 02-NOV-1999;
Location/Qualifiers
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Siciliano, M.J. and Liu, P.
In-situ hybridization probes for identification and banding of specific human chromosomes and regions
Patent: US 5538869-A 2 23-UTL-1996;
Location/Qualifiers
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AR083936
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Sequence 2 from patent US 5538869.
123816
123816.1 GI:1603686
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123816/c
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Homo sapiens (human)
Homo sapiens (human)
Homo sapiens (human)
Homo sapiens (human)

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

22 I (bases 1 to 19)

23 Tavtigian,S.V., Teng,D.H.F., Simard,J. and Rommens,J.M.
Chromosome 17q-1inked prostate cancer susceptibility gene
Chromosome 17q-1inked prostate cancer susceptibility gene
Arent: JP 2002529065-A,99 Po 10-SEP-2002;
MYRIAD GENETICS: INC, THE HOSPITAL FOR SICK CHILDREN
OS Homo sapiens (human)
PN JP 2002529065-A/99
PD 10-SEP-2002
PF 05-NOV-1999 US 60/107468
PR 06-NOV-1998 US 60/107468
PI SERAN V TAVTIGIAN, DAVID H F TENG, JACQUES SIMARD, JOHANNA M PI
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CO7K14/47,CO7K16/18,CO7K16/44,C12N1/15,C12N1/19,C12N1/21,C12N5/ PC
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A61P35/00,
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GOIN37/00,C12N15/00,A61K37/02,C12N5/00
Ghromosome 17q-linked prostate cancer susceptibility gene FH
Location/Qualifiers
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Chromosome 17q-linked prostate cancer susceptibility gene.
BD231547
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          linear
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0.7%; Score 17.6; DB 1;
Best Local Similarity 94.4%; Pred. No. 9.8e+02;
Matches 17; Conservative 1; Mismatches 0;
          DNA
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Gilliam, T. Conrad. and Tanzi, R. B.
Wilson's disease gene
Watent: US 5578493-A 2 26-NOV-1996;
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Sequence 2 from patent US 5578493.
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/organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"
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/organism="unknown"
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Detera-Wadleigh, S.D., Yoshikawa, T., Sanders, A.R. and Esterling, L.E. Polynucleotides encoding IMP.18p myo-inositol monophosphatase and methods of detecting said polynucleotides
Patent: US 6458532-A 86 01-00T-2002;
Location/Qualifiers
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Unclassified.
Unclassified.
I (bases 1 to 19)
Weber, J.L.
Length Jolymorphisms in (dC-dA).sub.n.(dG-dT).sub.n sequences and method of using the same
method of using the same
Location/Qualifiers
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0.7%; Score 17.4; DB 1; Length 19; 94.7%; Pred. No. 1e+03; Live 0; Mismatches 1; Indels
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0.7%; Score 17.4; DB 1; Length 19;
Best Local Similarity 94.7%; Pred. No. 1e+03;
Matches 18; Conservative 0; Mismatches 1; Indels
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Sequence 330 from patent US 5582979.
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Seguence 86 from patent US 6458532.
AR233457
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AR482557
AR482557.1 GI:47245080

    1. .19
/organism="unknown"
/wol_type="genomic DNA"

                                                                2280 CACCGTGTTAGCCAGGATG 2298
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131418.1 GI:1822209
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Best Local Similarity
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

1 (bases 1 to 19)
Landers,J.E., Jordan,B., Housman,D.E. and Charest,A.
Methods and products related to genotyping and DNA analysis
Patent: JP 2002525127-A 3 13-AUG-2002;
MASSACHUSETTS INSTITUTE OF TECHNOLOGY
OS Homo sapiens (human)
PN JP 2002525127-A/3
PP 24-SEP-1999 JP 2000572407
PR 25-SEP-1999 US 60/101757
PR 25-SEP-1999 US 60/101757
PR 25-SEP-1999 US 60/101757
PR 25-SEP-1999 US 60/10133/59,G01N33/59,G01N33/59,G01N37/00, PC
                                                                                                                         BD241056

19 bp DNA linear PAT 17-JUL-2003 Methods and products related to genotyping and DNA analysis. BD241056.1 GI:33050826
JP 2002525127-A/3. Homo sapiens (human)
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C12N15/00
Methods and products related to genotyping and DNA analysis FH
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Reichwald, K.
Morti variant associated with human obesity
Patent: WO 2003104489-A 107 18-DEC-2003;
Philipps-Universitaet Marburg (DE)
Location/Qualifiers
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/organism="synthetic construct"
/mol_type="unassigned DNA"
/mol_type="taxon:32630"
/noce="primer B6r"
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    .19
    /organism="Homo sapiens"
/mol_type="genomic DNA"
/db_xref="taxon:9606"

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                            CCTCAGCCTCCCAAATAGC 1
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             CCTCAGCCTCCCAATTAGC
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2175 CGGGTTCGCACCATTCTCC 2193
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their use in diagnostic
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                                                          1 (bases 1 to 19)
Landers, J., Jordan, B., Housman, D.E. and Charest, A.
Methods and productes related to genotyping and DNA analysis
Patent: US 670328-A 3 09-MAR-2004;
Location/Qualifiers
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Patent: WO 0109183-A 12 08-FEB-2001;
EPIDAUROS AG Biotechnologie Aktiengesellschaft (DE)
Location/Qualifiers
                                                                                                                                                                                                  0.7%; Score 17.4; DB 1; Length 19; 94.7%; Pred. No. 1e+03; tive 0; Mismatches 1; Indels
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94.7%; Pred. No. 1e+03;
Lve 0; Mismatches 1; Indels
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and therapeutic applications
Extent: WO 0109183-A 210 08-FEB-2001;
EPIDAUROS AG Biotechnologie Aktiengesellschaft
Location/Qualifiers
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/organism="synthetic construct"
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Sequence 212 from Patent WO0109183.
AX081968
                                                                                                                                                                    /mol_type="genomic DNA"
                                                                                                                                         1. .19
/organism="unknown"
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Matches 18; Conservative
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Best Local Similarity 94.7
Matches 18; Conservative
                               Unknown.
Unclassified.
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Polymorphisms in the human mdr-1 gene and their use in diagnostic
and therapeutic applications
Patent: WO 0109183-A 216 08-FEB-2001;
PRIDAUROS AG Blotechnologie Aktiengesellschaft (DE)
Location/Qualifiers
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their use in diagnostic
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                                                                                          0.7%; Score 17.4; DB 1; Length 19;
llarity 94.7%; Pred. No. 1e+03;
Conservative 0; Mismatches 1; Indels
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/note="synthetic"
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/organism="synthetic construct"
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Sequence 218 from Patent W00109183.
AX081974
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Sequence 216 from Patent WO0109183
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PAT 10-SEP-2001

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PAT 27-AUG-2002

LOCUS DEFINITION

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RESULT 886 AX116142

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AX081978/c LOCUS DEFINITION

RESULT 884

ORGANISM ACCESSION VERSION KEYWORDS SOURCE

AUTHORS REFERENCE

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vmglom gene and its mutations causing disorders with a vascular component
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A method of arraying genome clone
Patent: JP 2001321190-A 943 20-NOV-2001;
THE INSTITUTE OF PHYSICAL AND CHEMICAL RESEARCH, YUGENKAISHA
                                                                                                                                Genotyping reagents, kits and methods of use thereof
Patent: WO 0129262-A 1265 26-APR-2001;
Orchid BioSciences, Inc. (US)
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Pred. No. 1e+03;
0; Mismatches 1; Indels
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Pred. No. 1e+03;
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UNIVERSITE CATHOLIQUE DE LOUVAIN (BE)
Location/Qualifiers
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Polymorphisms in the human mdr-1 gene and their use in diagnostic
and therapeutic applications
Patent: WO 0109183-A 224 08-FEB-2001;
EPIDAUROS AG Biotechnologie Aktiengesellschaft (DE)
Location/Qualifiers
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their use in diagnostic
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Pred. No. le+03;
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Seguence 1265 from Patent WO0129262.
AX116142
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Sequence 222 from Patent WO0109183.
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REFERENCE AUTHORS TITLE

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chromosome 1p35-p36
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                             at 1p36.
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AB068733.1 GI:15129537
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C12N15/09,C12N15/09,C12M1/00,C12Q1/68,G01N33/53,G01N33/566, PC
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N JP 2001321190-A/2316
D 20-NOV-2001
F 12-MAR-2001 JP 2001068285
I B EICHI SOEDA
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Patent: JP 2001321190-A 2316 20-NOV-2001;
THE INSTITUTE OF PHYSICAL AND CHEMICAL RESEARCH, YUGENKAISHA
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/organism='Artificial Sequence'.
Location/Qualifiers
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/organism='Artificial Sequence'
Location/Qualifiers
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    .19
        /organism="synthetic construct"
/mol_type="genomic DNA"
        /db_xref="taxon:32630"

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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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JP 2001321190-A/2316.
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RESULT 890 AB068733/c

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AB068733 19 bp DNA linear SYN 21-MAY-2003
Synthetic construct DNA, reverse primer for human STS sts-D1S2728
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                                                                                                                                                                                                                                                                                                     Chen,Y.Z., Hayashi,Y., Wu,J.G., Takaoka,E., Maekawa,K.,
Watanabe,N., Inazawa,J., Hosoda,F., Arai,Y., Mizushima,H.,
Morohashi,A., Ohira,M., Nakagawara,A., Liu,S., Hoshi,M., Horii,A.
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/mol type="genomic DNA"
/mol type="genomic DNA"
1. 13
/note="reverse primer for human STS sts-D182728 at 1p36
sts-D182728 obtained from clones B351N1, B26G13, B26E12,
B39F12, Human BAC library RPCI-11"
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Submitted (04-AUG-2001) Akira Horii, Tohoku University School of
Submitted (04-AUG-2001) Akira Horii, Tohoku University School of
Medicine, Molecular Pathology; 2-1 Seiryomachi, Aoba-ku, Sendai,
Miyagi 980-8575, Japan (E-mail:horii@mail.cc.tohoku.ac.jp,
Tel:81-22-717-8042, Fax:81-22-717-8047)
Location/Qualifiers
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A BAC-based STS-content map spanning a 35-Mb region of human
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Best Local Similarity 94.7%; Pred. No. 1e+03;
Matches 18; Conservative 0; Mismatches 1; Indels
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Sequence 351 from patent US 5582979.
131439 1 GI:1822230
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/mol_type="unassigned DNA"
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Patent: US 6545137-A 286 08-APR-2003;
                        1. .20
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          Location/Qualifiers
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Conservative
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Unclassified.
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AR309436
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AX184102/c
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Todd,J.A., Hess,J.W., Caskey,C.T., Cox,R.D., Gerhold,D.,
Hammond,H., Hey,P., Kawaguchi,Y., Merziman,T.R., Metzker,M.L.,
Nakagawa,Y., Phillips,M.S. and Twells,R.C.J.
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AS Bennett,C.F., Freier,S.M. and Watt,A.T.
Antisense modulation of phospholipase A2, group VI (Ca2+-independent) expression
(Ca2+-independent) expression
NAL Patent: US 6410325-A 18 25-UNN-2002;
Location/Qualifiers
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Patent: US 6503152-A 95 07-JAN-2003;
Location/Qualifiers
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                             Sequence 18 from patent US 6410325.
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Sequence 95 from patent US 6503152.
AR271152. GI:29702455
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1 (bases 1 to 20)
Pelz,D.T.
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Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Todd,J.A., Hess,J.W., Caskey,C.T., Cox,R.D., Gerhold,D.,
Hammond,H., Hey,P., Kawaguchi,Y., Merriman,T.R., Metzker,M.L.,
Nakagawa,Y., Phillips,M.S. and Twells,R.C.J.
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Patent: WO 0142511-A 1855 14-JUN-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH (US); Ellipsis Biotherapeutics Corporation (CA)
Location/Qualifiers
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ch 0.7%; Score 17.4; DB 1; Length 20; Il Similarity 94.7%; Pred. No. 1e+03; 18; Conservative 0; Mismatches 1; Indels
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Location/Qualifiers
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Sequence 286 from patent US 6555654.
AR309436.1 GI:31701441
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Seguence 1855 from Patent WO0142511.
AX184102
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/db_xref="taxon:9606"
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/organism="unknown"
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PAT 18-SEP-2002

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PN JP 2002501376-A/258
PD 15-JAN-2002
PF 15-APR-1998 JP 1998543635
PR 15-APR-1999 US 60/043553,05-JUN-1997 US 60/048740 PI
JOHN ANDREW TODD, JOHN WILFRED HESS, CHARLES
THOWAS CASKEY, ROGER
PI DAVID GERHOLD, HOLLY HAMMOND, PATRICIA HEY
PC C12015/11,2, C12201/68,C07K14/705,C07K16/28,A61K38/17,PC
PC A61K49/00
CC Strandedness: Single;
CC Topology: Linear;
FH Key
                                                                                                                                                                                                                                                                                                            Novel LDL-receptor
Patent: JP 2002501376-A 258 15-JAN-2002;
THE WELLCOME TRUST LTD AS TRUSTEE TO THE WELLCOME TRUST, MERCK & CO
INC
                                                                                                                                                                                                                                      Chlamydia sp.
Chlamydia sp.
Bacteria; Chlamydiae; Chlamydiales; Chlamydia.
1 (bases 1 to 20)
Todd,J.A., Hess,J.W., Caskey,C.T., Cox,R.D., Gerhold,D., Hammond,H.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1 (bases 1 to 21)
Fu,Y.-H., Yu,C.-E., Oshima,J., Mulligan,J.T. and Schellenberg,G.D. Gene products related to werner's syndrome
Patent: US 6583112-A 28 24-JUN-2003;
Location/Qualifiers
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Sequence 28 from patent US 6583112.
AR345147

    .20
    /organism="Chlamydia sp./mol_type="genomic DNA" /db_xref="taxon:35827"

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/organism="unknown"
/mol_type="genomic DNA"
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JP 2002501376-A/258.
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                                        20 CTCAGCCTCCCAAGTAGCT
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Best Local Similarity
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BD106243
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                                                                                                                                                                                                                                                    van Roy, F., Vanlandschoot, A. and Janssens, B.

Novel cdnas encoding catenin-binding proteins with function in signalling and/or gene regulation
Patent: WO 147954-A 30 05-JUL-2001;
Vlaams Interuniversitair Instituut voor Biotechnologie vzw. (BE)
Location/Qualifiers
1..20

/organism="synthetic_construct"
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JP 2001321190-A/1556
JP 2001321190-A/1556
12-MAR-2001
EIICH SOEDA
C12N15/09,C12N15/09,C12M1/00,C12Q1/68,G01N33/53,G01N33/566,
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Patent: JP 2001321190-A 1556 20-NOV-2001;
THE INSTITUTE OF PHYSICAL AND CHEMICAL RESEARCH, YUGENKAISHA
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0.7%; Score 17.4; DB 1; Length 20;
Best Local Similarity 94.7%; Pred. No. 1e+03;
Matches 18; Conservative 0; Mismatches 1; Indels
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'organism='Artificial Sequence'
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/db_xref="taxon:32630"
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Sequence 30 from Patent W00147954.
AX188411
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JP 2001321190-A/1556.
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synthetic construct

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artificial sequences.

I (bases 1 to 22)

Sethor, O., Ichiro, N. and Hiroshi, Y.

Method for distinguishing eucaryctic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences

and primer to be used therein

IL Patent: JP 200023571-A 13 25-JAN-2000;

NATIONAL RESEARCH INSTITUTE OF AQUACULTURE

OS Artificial Sequence

PN JP 200023571-A/13

PD 25-JAN-2000

PF 10-JUL-1998 JP 1998195692

PR 10-JUL-1998 JP 1998195692
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Synthetic construct
Synthetic construct
artificial sequences.

E 1 (bases 1 to 22)
Sethoro, O., Ichiro, N. and Hiroshi, Y.

Method for distinguishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein

I. Patent: JP 2000033671-A 14 25-JAN-2000;
NATIONAL RESEARCH INSTITUTE OF AQUACULTURE
OS Artificial Sequence
PN JP 2000023671-A/14
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Method for distinguishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein.

E31641.1 GI:13018551
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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                                            2294 GGATGGTCTCGATCTCCTG 2312
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                                                                     GGATGGTCTCGAACTCCTG 3
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Conservative
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18;
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synthetic construct
artificial sequences.

E 1 (bases 1 to 22)

Is in (bases 1 to 22)

Method for distinguishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein

In Patent: JP 2000021671-A 15 25-JAN-2000;

NATIONAL RESEARCH INSTITUTE OF AQUACULTURE
OS Artificial Sequence
PN JP 2000021871-A/15

PP 10-JUL-1998 JP 1998195692
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Method for distinguishing eucaryotic individual based on PCR finger
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                                                                                                              ce 1. .22
/organism='Artificial Sequence'.
Location/Qualifiers
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/organism='Artificial Sequence'.
Location/Qualifiers
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C12N15/09, C12Q1/68, C12N15/00
ICHIRO OHARA, ICHIRO NAKAYAMA, HIROSHI YASUE
C12N15/09, C12Q1/68, C12N15/00
                                                                                      Location/Qualifiers
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/organism="synthetic construct"
/mol_type="genomic DRA"
/db_xref="taxon:32630"
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Matches 18; Conservative
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Wed Jan 26 12:47:31 2005

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JP 2000023671-A/18.
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sm synthetic construct
sm stificial sequences.

1 (bases 1 to 22)

1 (bases 1 to 22)

2 Ichiro, N. and Hiroshi, Y.

2 Ichiro, O., Ichiro, N. and Hiroshi, Y.

2 Ichiro, O., Ichiro, N. and Hiroshi, Y.

3 Ichiro, O., Ichiro, N. and Hiroshi, Y.

3 Ichiro, O., Ichiro, N. and Hiroshi, Y.

4 Ichiro, O., Ichiro, N. and Hiroshi, Y.

5 Ichiro, O., Ichiro, N. and Hiroshi, Yasue

5 Ichiro, O., Ichiro, N. and Hiroshi, Yasue

6 Ichiro, O., Ichiro, O., Ichiro, Qualifiers

7 Ichiro, O., Ichiro, Ichiro, O., Ichiro, Qualifiers

8 Ichiro, O., Ichiro, Ichiro, Ichiro, Qualifiers

8 Ichiro, O., Ichiro, N. and Hiroshi, Yasue

8 Ichiro, O., Ichiro, Ichiro, Ichiro, Qualifiers

8 Ichiro, O., Ichiro, Ichiro, Ichiro, Qualifiers

8 Ichiro, O., Ichiro, Ichiro, Ichiro, Ichiro, Qualifiers

8 Ichiro, O., Ichiro, Ichiro,
                                                                                                                                                                                                        artificial sequences.

1 (Dases I to 22)
Ichiro,O., Ichiro,N. and Hiroshi,Y.
Method for distinguishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein.

Patent: JP 2000023571-A 16 25-JNN-2000;
NATIONAL RESEARCH INSTITUTE OF AQUACULTURE
OS Artificial Sequence
PN JP 2000023671-A/16
PD 25-JAN-2000
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print with the use of restriction primer of inter-SINE sequences and primer to be used therein. \mathbf{E31643}
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Location/Qualifiers
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C12N15/09,C12Q1/68,C12N15/00
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/organism="synthetic construct"
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/db_xref="taxon:32630"
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JP 2000023671-A/16.
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synthetic construct
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E31644.1 GI:13018554
JP 2000023671-A/17.
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synthetic construct
synthetic construct
artificial sequences.

1 (bases 1 to 22)
Ichiro,O., Ichiro,N. and Hiroshi,Y.
Method for distinguishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein
Patent: JP 2000023671.A 18 25-JAN-2000;
NATIONAL RESEARCH INSTITUTE OF AQUACULTURE
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Method for distinguishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein.

E31645.1 GI:13018555
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artificial sequences.

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Ichiro, O., Ichiro, N. and Hiroshi, Y.
Method for distinguishing eucaryotic individual based on PCR finger
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llarity 94.7%; Pred. No. 1e+03;
Conservative 0; Mismatches 1; Indels

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/organism='Artificial Sequence'

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C12N15/09, C12Q1/68, C12N15/00
/organism="synthetic construct"
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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JP 2000023671-A/19.
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artificial sequences.

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E 1 (bases 1 to 22)

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Method for distinguishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein.
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94.7%; Pred. No. 1e+03;
tive 0; Mismatches 1; Indels
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"

    .22
    ^2rganism="synthetic construct"
    ^mol_type="genomic DNA"
    ^db_xref="taxon:32630"

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JP 2000023671-A/20.
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synthetic construct
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/organism='Artificial Sequence'.
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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Pred. No. 1e+03;
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B31649
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FEATURES

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E31652

DNA linear PAT 18-JUN-2001

Print with the use of restriction primer of inter-SINE sequences and primer to be used therein.

E31652

DNA 1308562

SNA STATE SEQUENCES

SYNCHOLOGOS 13018562

SYNCHOLOGOS 14018562

SYNCHOLOGOS 14018562

SOURCE SOURCES

SYNCHOLOGOS 14018562

SYNCHOLOGOS 14018563

SYNCHOL
synthetic construct
synthetic construct
artificial sequences.

1 (bases 1 to 22)
Ichiro,O., Ichiro,N. and Hiroshi,Y.
Ichiro,O., Ichiro,O.
Ichiro,O.
Ichiro,O., Ichiro,O.
Ichir
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Location/Qualifiers
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
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O.7%; Score 1...,
Best Local Similarity 94.7%; Pred. No. 1e+0.
Matches 18; Conservative 0; Mismatches
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I (bases 1 to 22)

I (bases 2 to 22)

I (bases 3 to 22)

I (bases 3 to 22)

I (bases 2 to 22)

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We thod for distinguishing evearyotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein.

E31651.

E31651.

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JP 2000023671-A/24.
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    .22
    /organism='Artificial Sequence'.

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Best Local Similarity 94.7%; Pred. No. 1e+03;
Matches 18; Conservative 0; Mismathham
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/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                 Location/Qualifiers

    .22
    ^organism="synthetic construct"
/mol_type="genomic DNA"
    /db_xref="taxon:32630"

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E31650.1 GI:13018560
JP 2000023671-A/23.
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DEFINITION

ACCESSION VERSION KEYWORDS

RESULT 912

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ACCESSION VERSION KEYWORDS SOURCE ORGANISM

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1 (bases 1 to 2)
...dORS Ichiro,0. Ichiro,N. and Hiroshi,Y.
Method for distinguishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein.
JOURNAL Patent: JP 2000023671-A 26 25-JAN-2000;
NATIONAL RESEARCH INSTITUTE OF AQUACULTURE
OS Artificial Sequence
PN 10-JUL-1998 JP 1998195692
PR 10-JUL-1998 JP 1998195692
PR 10-JUL-1998 JP 1998195692
PR 10-JUL-1998 JP 1998195692
PR F 10-JUL-1998 JP 1998195692
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1 (bases 1 to 22)
Ichiro,0., Ichiro,N. and Hiroshi,Y.
Ichiro,0., Ichiro,N. and Hiroshi,Y.
Method for distringuishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein
Patent: JP 20000235671-A 27 25-JAN-2000;
NATIONAL RESEARCH INSTITUTE OF AQUACULTURE
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                  0; Mismatches
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1 (bases 1 to 22)
Ichiro,O., Ichiro,N. and Hiroshi,Y.
Method for distinguishing eucaryotic individual based on PCR finger
print with the use of restriction primer of inter-SINE sequences
and primer to be used therein.
Patent: JP 2000023671-A 28 25-JAN-2000;
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E31655
E31655.1 GI:13018565
JP 2000023671-A/28.
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OS Artificial Sequence
PN JP 2000023671-A/28
PD 25-JAN-2000
PF 10-JUL-1998 JP 1998195692
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PI ICHIRO OHARA, ICHIRO NAKAYAMA, HIROSHI YASI
PC C12N15/09, C12Q1/68, C12N15/00
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Pred. No. 1e+03;
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Best Local Similarity 94.7%;
Matches 18; Conservative
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Bynthetic construct

Synthetic construct

artificial sequences.

Is (bases 1 to 22)

S Ichiro, O., Ichiro, N. and Hiroshi, Y.

Method for distinguishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein

And primer to be used therein

NATIONAL RESEARCH INSTITUTE OF AQUACULTURE

OS Artificial Sequence

PU JP 2000023671-A/30
                                                                                            Synthetic construct
artificial sequences.

1 (bases 1 to 22)
1 (chiro, 0., Ichiro, N. and Hiroshi, Y.
Method for distinguishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein Patent: JP 200023671-A 29 25-JAN-2000;
NATIONAL RESEARCH INSTITUTE OF AQUACULTURE
OS Artificial Sequence
PN JP 2000023671-A/29
PD 25-JAN-2000
Method for distinguishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein.

E31656. E31656. E3161018566
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22 bp DNA linear PAT 18-JUN-2001 Method for distinguishing eucaryotic individual based on PCR finger print with the use of restriction primer of inter-SINE sequences and primer to be used therein.
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/organism='Artificial Sequence'
Location/Qualifiers
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synthetic construct
synthetic construct
artificial sequences.

I (bases 1 to 23)
Cotaki,T., Masuda,Y., Takatsu,Y., Watanabe,T., Terao,Y., Shintani,Y.
and Hinman,S.
Novel phistological active peptide and its use
Patent: WO 02062944-A 12 15-AUG-2002;
TAKEDA CHEMICAL INDUSTRIES LTD, TETSUYA OTAKI, YASUSHI WASUDA,
YOSHIHIRO TAKATSU,TAKUYA WATANABE, YASUKO TERAO, YASUSHI SHINTANI,
SHUJI HINUMA
OS Artificial Sequence
PP 15-AUG-2002
PP 02062944-A/12
PP 01-FEB-2001 UP 01P 026820
PR 02-FEB-2001 UP 01P 026820
PR 02-FEB-2001 UP 01P 026820
PR TETSUYA OTAKI, YASUSHI MASUDA, YOSHIHIRO TAKATSU, TAKUYA
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CO7KK4/47, CO7KI4/705, C12N15/12, C12P21/02, C07K16/18, A61K67/027,
C12N5/10,
G01N13/15, G01N13/50, A61P1/00
DNA primer, hBv8-WR primer
Key
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Pred. No. 1e+03;
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Sequence 4 from patent US 5597694.
134290.
134290.1 GI:1825081
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WO 02062944-A/12.
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AUTHORS TITLE

JOURNAL

RESULT 921 AR080244/c

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DEFINITION

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ACCESSION VERSION KEYWORDS

ORGANISM

SOURCE

AUTHORS TITLE

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PAT 12-JUN-2003
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Unknown.
Unknown.
Unclassified.
1 (bases 1 to 22)
1 (bases 1 to 22)
1 (bases 1 to 22)
Plevy, S. E., Rotter, J. I., Targan, S. R., Toyoda, H. and Yang, H.
Methods of screening for Crohn's disease using TNF microsatellite
allelse
Patent: US 6534263-A 1 18-MAR-2003;
Location/Qualifiers
                                                                                                                                                                                                             Unclassified.

1 (bases 1 to 22)
Plevy, S.E., Targan, S.R., Taylor, K. and Barry, M.J.
Methods of diagnosing clinical subtypes of crohn's disease with characteristic responsiveness to anti-Thl cytokine therapy
Patent: US 6183951-A 1 06-PEB-2001;
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1 (bases 1 to 22)
Taylor,K.D., Rotter,J.I. and Yang,H.
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Seguence 28 from patent US 6376176.
AR367700
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6183951.
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Sequence 1 from patent US 6534263.
AR287807
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Sequence 1 from patent US
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Unclassified.

(I (bases 1 to 22)
Plevy,S.E., Rotter,J.I., Targan,S.R., Toyoda,H. and Yang,H.
Methods of screening for Crohn's disease using TNF microsatellite alleles
Munroe, D.J. and Housman, D.B.
Interspersed repetitive element-bubble amplification of nucleic
acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Unclassified.

1 (bases 1 to 22)
Plevy, S.E. and Targan, S.R.
Methods of diagnosing a medically resistant clinical subtype of ulcerative colitis
Patent: US 5968741-A 1 19-OCT-1999;
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Best Local Similarity 86.4%; Pred. No. 1.1e+03;
Matches 19; Conservative 0; Mismatches 3;
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/organism="unknown"
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                                               Patent: US 5597694-A 4 28-JAN-1997;
Location/Qualifiers
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Location/Qualifiers
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Sequence 1 from patent US 5968741.
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Garini,Y., Cabib,D., Buckwald,R.A., Ried,T. and Soenksen,D.G.
Method for simultaneous detection of multiple fluorophores for in
situ hybridization and multicolor chromosome painting and banding
Patent: US 5817462-A 2 06-OCT-1998;
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Methods of using a major histocompatibility complex class III haplotype to diagnose Crohn's disease Patent: US 6376176.A. 28 27-APR-2002; Location/Qualifiers
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Methods of using a major histocompatibility complex class iii
haplotype to diagnose crohn's disease
Patent: WO 01200156.A 28 22-MAR.2001;
CEDARS-SINAI MEDICAL CENTER (US)
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Best Local Similarity 86.4%; Pred. No. 1.1e+03;
Matches 19; Conservative 0; Mismatches 3;
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    .22
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Sequence 28 from Patent WO0120036.
AX098591
AX098591.1 GI:13537855
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Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;

Mammalla; Eutheria; Primates; Catarrhini; Hominidae; Homo.

B Pavco, P.A., Roberts, E., Jarvis, T., Coeshott, C. and Mcswiggen, J.A.

Method and reagent for treating diseases or conditions concerning molecule participating in vasculogenic response

RIBOZYNE PHARMACETICALS INC

OS Homo sapiens (human)

PN JP 2002509721-A/6057

PP 24-MAR-1999 UB 60/079678

PR 27-MAR-1999 UB 60/079678

PR 27-MAR-1999 US 60/079678

PI PAMELA A PAVCO, ELISABETH ROBERTS, THALE JARVIS, CLAIRE COESHOTT, AND A PANCO A PANCO AND A PANCO A PANCO AND A PANCO AND A PANCO A PANC
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24-MAR-1999 UP 2000541291
27-MAR-1998 US 60/079678
PAMELA A PAVCO, ELISABETH ROBERTS, THALE JARVIS, CLAIRE COESHOTT, JAMES A MCSWIGGEN
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PC
C12N15/09,A61K31/7088,A61K31/7125,A61K48/00,A61P3/10,A61P17/06, PC
Method and reagent for treating diseases or conditions concerning molecule participating in vasculogenic response.

BD203031
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Method and reagent for treating diseases or conditions concerning molecule participating in vasculogenic response.

BD203159
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1 (Dassel 1 to 17)
Pavco, P.A., Roberts, E., Jarvis, T., Coeshott, C. and Mcswiggen, J.A. Method and reagent for treating diseases or conditions concerning molecule participating in vasculogenic response Patent: JP 2002509721-A 6185 02-APR-2002;
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PC AG1P35/00,AG1P43/00,C12N5/10,C12N9/00//AG1K35/76,C12N15/00,
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participating in vasculogenic response
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Pred. No. 1e+03;
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/organism="Homo sapiens"
/mol_type="genomic RNA"
/db_xref="taxon:9606"
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JP 2002509721-A/6185.
Homo sapiens (human)
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Telerman,A., Amson,R. and Tuijnder,M. Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses and their use as medicines
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Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or resistance to viruses and their use as medicines
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       Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Molecular Engines Laboratories (FR)
Location/Qualifiers
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Seguence 2065 from Patent WO03004526.
AX673620
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Molecular Engines Laboratories (FR)
Location/Qualifiers
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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/mol_type="unassigned DNA"
/db_xref="taxon:9606"
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PC A61P35/00,A61P43/00,C12N5/10,C12N9/00//A61K35/76,C12N15/00, PC
RIBOZYME PHARMACEUTICALS INC
OS Homo gapiens (human)
PN 07 2002509721-A/6185
PD 02-APR-2002
PF 24-MAR-1999 JP 2000541291
PR 27-MAR-1999 US 60/079678
PI JAMES A PAVCO, ELISABETH ROBERTS, THALE JARVIS, CLAIRE COESHOTT, PI JAMES A MCSWIGGEN
PI JAMES A MCSWIGGEN
CLAIS, 09, A61K31/7088, A61K31/7125, A61K48/00, A61P3/10, A61P17/06, PC
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100.0%; Pred. No. 1e+03;
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AX671897
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Disease-associated gene
Datent: WO 0206312-A 75 24-JAN-2002;
Location/Qualifiers
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AX393527
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/organism="Homo sapiens"
/mol_type="genomic RNA"
/db_xref="taxon:9606"
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
mdz12
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0.7%; Score 17; DB 1; Length 17;
Best Local Similarity 100.0%; Pred. No. 1e+03;
Matches 17; Conservative 0; Mismatches 0; Indels
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Patent: WO 03004526-A 3149 16-JAN-2003;
Molecular Engines Laboratories (FR)
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Aeomica, Inc. (US)
Location/Qualifiers
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Sequence 5449 from Patent EP1281758.
AX692717
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Sequence 5448 from Patent EP1281758.
AX692716.1 GI:29415674
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    /organism="Homo sapiens"
/mol_type="unassigned DNA"
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AX692716
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Sequence 5451 from Patent BP1281758.
AX692719
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Aeomica, Inc. (US)
Location/Qualifiers
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Sequence 5450 from Patent EP1281758.
AX692718
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Location/Qualifiers
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RESULT 938

0.7%; Score 17; DB 1; Length 17;

Query Match

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Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo
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Sequence 5456 from Patent
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and mdz12
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Homo sapiens
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
                                                                                                                   Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                            Shannon,M., Gu,Y. and Nguyen,C.T.
Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
mdz12
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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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             AX692720 17 bp DNA Sequence 5452 from Patent EP1281758. AX692720
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Shannon, M., Gu, Y. and Nguyen, C.T.
Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
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Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
mdz12
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Acomica, Inc. (US)
Location/Qualifiers
                                                                   Patent: BP 1281758-A 5454 05-FEB-2003;
Aeomica, Inc. (US)
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Homo sapiens
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
mdz12
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                        Sequence 5459 from Patent EP1281758. AX692727
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AX692728 1GI:29415686
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Sequence 5461 from Patent BP1281758.
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
mdz12
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo
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Aeomica, Inc. (US)
Location/Qualifiers
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Patent: BP 1281758-A 5458 05-FBB-2003;
Aeomica, Inc. (US)
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Sequence 5458 from Patent EP1281758.
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AX692726.1 GI:29415684
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Location/Qualifiers
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Sequence 5463 from Patent EP1281758.
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Sequence 5462 from Patent EP1281758.
AX692730

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AX692729.1 GI:29415687
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Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
mdz12
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Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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100.0%; Pred. No. 1e+03;
tive 0; Mismatches 0; Indels
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Patent: EP 1281758-A 5463 05-FEB-2003;
Acomica, Inc. (US)
Location/Qualifiers
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Aeomica, Inc. (US)
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Aeomica, Inc. (US)
Location/Qualifiers
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Sequence 5464 from Patent EP1281758.
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Sequence 5465 from Patent EP1281758.
AX692733 1 GI:29415691
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Best Local Similarity 100.0
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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Acomica, Inc. (US)
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Sequence 5468 from Patent EP1281758.
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and mdz12
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Aeomica, Inc. (US)
Location/Qualifiers
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Sequence 5466 from Patent EP1281758.
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Sequence 5467 from Patent EP1281758.

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Bukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Sequences involved in phenomena of tumour suppression, tumour reversion, apoptosis and/or virus resistance and their use as medicines
Patent: WO 03025175-A 5752 27-MAR-2003;
Molecular Engines Laboratories (FR)
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Molecular Engines Laboratories (FR)
Location/Qualifiers
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                               Shannon, M., Gu, Y. and Nguyen, C.T.
Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
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Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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Aeomica, Inc. (US)
Location/Qualifiers
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Aeomica, Inc. (US)
Location/Qualifiers
                                                                                                                                 Patent: EP 1281758-A 5470 05-FEB-2003,
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Sequence 5471 from Patent EP1281759.
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Location/Qualifiers
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AX692739.1 GI:29415697
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DEFINITION ACCESSION VERSION

RESULT 957 AX692739

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Methods, kits and compositions pertaining to the suppression of
detectable probe binding to randomly distributed repeat sequences
in genomic nucleic acid
Patent: WO 03027328-A 10 03-APR-2003
Boston Probes, Inc. (US); DakoCytomation Denmark A/S (DK)
Location/Qualifiers
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Methods, kits and compositions pertaining to the suppression of
detectable probe binding to randomly distributed repeat sequences
in genomic nucleic acid
Patent: WO 03027328-A 22 03-APR-2003;
Boston Probes, Inc. (US); DakoCytomation Denmark A/S (DK)
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Sequence 22 from Patent WO03027328.
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Polymorphisms in the human mdr-1 gene and their use in diagnostic
and therapeutic applications
Patent: WO 0109183-A 211 08-FEB-2001;
EPIDAUROS AG Blotechnlogie Aktiengesellschaft (DE)
Location/Qualifiers
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their use in diagnostic
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Sequence 211 from Patent W00109183. AX081967
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Sequence 213 from Patent WO0109183.
AX081969
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Matches 17; Conservative
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Best Local Similarity 89.5%;
Matches 17; Conservative
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                                                          Brinkmann, U., Hoffmeyer, S., Eichelbaum, M. and Roots, I.
Polymorphisms in the human mdr-1 gene and their use in diagnostic
and therapeutic applications
Patent: WO 0109183-A 217 08-FEB-2001;
EPIDAUROS AG Biotechnologie Aktiengesellschaft (DE)
Location/Qualifiers
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Polymorphisms in the human mdr-1 gene and their use in diagnostic
and therapeutic applications
Patent: WO 0109183-A 219 08-FEB-2001,
EPIDAUROS AG Biotechnologie Aktiengesellschaft (DE)
Location/Qualifiers
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/db_xref="taxon:32630"
/note="y=t or c"
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0.7%; Score 17; DB 1;
Best Local Similarity 89.5%; Pred. No. 1e+03;
Matches 17; Conservative 1; Mismatches
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0.7%; Score 17; DB 1;
Best Local Similarity 89.5%; Pred. No. 1e+03;
Matches 17; Conservative 1; Mismatches
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AX081979.1 GI:13170786
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                synthetic construct artificial sequences.
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Heinrich, G. and Kerb, R. Methods for the treatment of cancer with irinotecan based on CYP3A5 Patent: WO 03013534-A 523 20-FEB-2003; Epidauros Biotechnologie AG (DE)
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Polymorphisms in the human mdr-1 gene and their use in diagnostic
and therapeutic applications
Patent: WO 010913-A 225 08-FEB-2001;
EPIDAUROS AG Biotechnologie Aktiengesellschaft (DE)
Location/Qualifiers
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Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Best Local Similarity 89.5%; Pred. No. 1e+03;
Matches 17; Conservative 1; Mismatches
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Seguence 523 from Patent WO03013534.
AX706826
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Sequence 225 from Patent WO0109183.
AX081981 GI:13170788
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Methods for the treatment of cancer with irinotecan based on CYP3A5
Patent: WO 03013334-A 524 20-FEB-2003;
Epidauros Biotechnologie AG (DE)
Location/Qualifiers
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Methods for the treatment of cancer with irinotecan based on CYP3A5
Patent: WO 03013534-A 527 20-FEB-2003;
Epidauros Biotechnologie AG (DE)
Location/Qualifiers
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Mammalla, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Seguence 524 from Patent WO03013534.
AX706827
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    1.19
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             17; Conservative
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Heinrich, G. and Kerb, R.
Methods for the treatment of cancer with irinotecan based on CYP3A5
Patent: WO 03013534-A 528 20-FEB-2003;
Epidauros Biotechnologie AG (DE)
Location/Qualifiers
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Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Sequence 524 from Patent W003013536.
AX707757.1 GI:29563930
Sequence 528 from Patent WO03013534. AX706831
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Sequence 523 from Patent WO03013536.
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AX707756.1 GI:29563929
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Homo sapiens
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0.7%; Score 17; DB 1; Length 20;
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                                             Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Buteleostomi,
Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
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Methods for treatment of cancer using irinotecan based on UGT1A1
Patent: WO 03101536-A 527 20-FEB-2003;
Epidauros Biotechnologie AG (DE)
Location/Qualifiers
                                                                                                       Methods for treatment of cancer using irinotecan based on UGT1A1 Patent: WO 03013536-A 524 20-PEB-2003; Epidauros Biotechnologie AG (DE) Location/Qualifiers
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llarity 89.5%; Pred. No. 1e+03;
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Sequence 527 from Patent WO03013536.
AX707760
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AX707761
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/organism="Homo sapiens"
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                Homo sapiens (human)
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//db_xref="taxon:32630"
//db_xref="based on Homo sapiens. Each nucleobase may be part
of a ribonucleotide, deoxyribonucleotide, or nucleotide
analog-n = r or U"
Methods for treatment of cancer using irinotecan based on UGT1A1 Patent: WO 03013536-A 528 20-FEB-2003; Epidauros Biotechmologie AG (DE)
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Antisense iap nucleobase oligomers and uses thereof
Patent: WO 03080638-A 164 02-OCT-2003;
Aegera Therapeutics Inc. (CA)
Location/Qualifiers
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1 (bases 1 to 20)
Sidransky,D.
Detection of neoplasia by analysis of saliva
Patent: US 6235470-A 155 22-MAY-2001;
Location/Qualifiers
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Sequence 164 from Patent WO03080638.
AX923729
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Sequence 155 from patent US 6235470.
AR152875
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AX526494.1 GI:25171301
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Unknown.
Unclassified.

E 1 (bases 1 to 20)

S Zhang,H. and Cowsert,L.M.
Antisense modulation of caspase 8 expression

NAL Patent: US 6258600-A 94 10-JUL-2001;

Location/Qualifiers

Location/Qualifiers
100.0%; Pred. No. 1.1e+03;
tive 0; Mismatches 0;
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/db_xref="texon:32630"
/note="Synthetic Primer"
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Sequence 94 from patent US 6258600.
AR162414
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AX477118
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Sequence 209 from Patent W00220847.
AX526494
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AR162414/c
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Location/Qualifiers
                                                                                                     Bodnar,J.S., Castellani,L.W., Chatterjee,A., de Jong,P.,
Lusis,A.J., Ohmen,J., Ross,D., Tafuri,S. and Wu,C.
Gene and sequence variation associated with lipid disorder
Patent: WO 0220847-A 209 14-MAR-2002;
THE REGENTS OF THE UNIVERSITY OF CALIFORNIA (US)
Location/Qualifiers
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synthetic construct
artificial sequences.
1 (bases 1 to 20)
Sidlanski, D.
Detection of neoplasia by analysis of saliva
Patent: JP 2002505888-A 155 26-FEB-2002;
THE JOHNS HOPKINS UNIVERSITY SCHOOL OF MEDICINE
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Detection of neoplasia by analysis of saliva.
BD134331
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/mol_type="unaasigned DNA"
/mol_txref="taxon:32630"
/nore="Synthetic Primer"
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Sequence 75 from Patent WO03062469.
AX811386
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JP 2002505888-A/155.
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synthetic construct
artificial sequences.
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artificial sequences.
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2114 CICTGTTACCCAGGCTGGAGTGCAGTGGTGATCTTGGCTCA
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/organism="unknown"
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AR124511/c
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Genetic and physical mapping of simple sequence repeat containing sequence tagged sites from the human genome
Unpublished (1994)
Original source text: Home sapiens DNA.
Submitted by: Utah Center for Human Genome Research University of Utah, Dept. of Human Genetics
2160 Bocles Institute of Human Genetics
Salt Lake City, UT 84112
e-mail: sts@corona.med.utah.edu
Primer A: CACTGCACTCCAGGCTGGG
Primer B: AGGTTGAGGCTGCAGTGAGC
End to Label: Primer B
PCR Profile:
Initial Denaturation: 94C 3008ec
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Human STS UT5223, 5' primer bind, sequence tagged site.
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94 C 10 sec.
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STS; PCR primer; STS sequence; dinucleotide repeat; microsatellite
NAT, microsatellite marker; sequence tagged site.
Homo sapiens (human)
Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi; Mammalia, Butheria, Primates, Catarrhini, Hominidae, Homo.
I tosas I to 64.
Gerken, S.C., Matsunami, N., Plaetke, R., Albertsen, H., Ballard, L., Melis, R., Lawrence, E., Moore, M., Holik, P.R., Carlson, M., Zhao, X., Robertson, M., Bradley, P., Elsner, T., Tingey, A., Lalouel, J.-M. and
                                                                                                                                                                                                                                                                                                                           Gaps
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                                                                                                                                                ce 1. .20 /organism='Artificial Sequence'. Location/Qualifiers
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    .20
    ^coganism="synthetic construct"
    ^mol type="genomic DNA"
    ^db_xref="taxon:32630"

                                                                                                                                      Location/Qualifiers
                                             10-MAR-1999 JP 2000535774
10-MAR-1998 US 09/038637
DAVID SIDLANSKI 05/00
C12N15/09, C12Q1/68, C12N15/00
nucleotide
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/organism="Homo sapiens"
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/db_xref="taxon:9606"
5. -723
/evidence=experimental
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Artificial Sequence
JP 2002505888-A/155
26-FBB-2002
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Best Local Similarity 61.99
Matches 26; Conservative
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PAT 07-SEP-2000
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Unknown.
Unclassified.

1 (bases 1 to 20)

1 (bases 1 to 20)

Antisense oligonucleotide compositions and methods for the inibition of c-Jun and c-Fos
Patent: US 598558-A 25 16-NOV-1999;

Location/Qualifiers
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( Chases 1 to 20)

Monia, B.P. and Cowsert, L.M.

Antisense modulation of telomeric repeat binding factor 1
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                              5 CACTGCACTCCAGGCTGGGANCAGAGCAAGACTCTGTCTCA 46
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0.7%; Score 16.8; DB 1;
Best Local Similarity 90.0%; Pred. No. 1.1e+03;
Matches 18; Conservative 0; Mismatches 2;
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Patent: US 6130088-A 38 10-OCT-2000;
Location/Qualifiers
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Sequence 80 from patent US 6171860.
AR124511
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Sequence 38 from patent US 6130088.
AR112674
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AR086204 GI:10012970
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Query Match
Best Local Similarity 90.0
Matches 18; Conservative
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AR176770
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AR162415/c
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Sidransky, D.
Detection of neoplasia by analysis of saliva
Patent: US 6235470-A 135 22-MAY-2001;
Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                           Unknown.
Unclassified.
I (bases 1 to 20)
S Baker, B.F. and Cowsert, L.M.
Antisense inhibition of rank expression
AL Patent: US 6171860-A 81 09-JAN-2001;
Location/Qualifiers
20
            1 (bases 1 to 20)
Baker, B.F. and Cowsert, L.M.
Antisense inhibition of rank expression
Patent: US 6171860-A 80 09-JAN-2001;
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Sequence 135 from patent US 6235470.
AR152855.
AR152855.1 GI:15120387
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/organism="unknown"
/mol_type="unassigned DNA"
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Sequence 81 from patent US 6171860.
AR124512 AR124512.1 GI:14109873
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/organism="unknown"
/mol_type="unassigned DNA"
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PAT 17-DEC-2001
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Unknown.
Unclassified.
Unclassified.
I (bases 1 to 20)
Miraglia,L.J., Nero,P., Graham,M.J. and Monia,B.P.
Antisense oligonucleotide modulation of human mdm2 expression
Patent: US 6239921-A 26 29-MAY-2001;
Location/Qualifiers
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Dean, N.M., McKay, R., Miraglia, L. and Baker, B.
Antisense oligonuclectide compositions and methods for the modulation of strivating protein 1
Patent: US 6312900-A 25 06-NOV-2001;
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Sequence 25 from patent US 6312900.
AR176770
AR176770.1 GI:17919125
AR154609 20 bp 1
Sequence 26 from patent US 6238921.
AR154609.1 GI:15122662
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Sequence 95 from patent US 6258600.
AR165415
AR162415.1 GI:16229593
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CQ771171.1 GI:45125304
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JP 200252013-A/5
06-AUG-2002
14-SEP-1999 JP 2000570197
16-SEP-1999 US 09/154444
PAUL O SHEPPARD, KEVIN P FOLEY
C12N15/09, A61K38/00, A61K39/395, A61P1/04, A61P1/14, A61P3/09,
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PC G01N33/50, G01N33/53, G01N33/577, C12N15/00, C12N5/00, A61K37/02
Oligonucleotide primer ZC12502
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Pred. No. 1.1e+03;
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/organism="synthetic construct"
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synthetic construct
artificial sequences.
1 (bases 1 to 20)
Sheppard,P.O. and Foley,K.P.
Sheppard, P.O. 2002524103-A 5 06-AUG-2002,
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               1. .20
/organism="unknown"
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Location/Qualifiers
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BD237996
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JP 2002524103-A/5.
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CQ758958/c
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Vlaams Interuniversitair Instituut voor Biotechnologie vz; w. (BE) Location/Qualifiers
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/organism="synthetic construct"
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/db xref="taxon:32630"
/noTe="Forward primer of exon 1, gene ZNF9"
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Gene assay method for predicting glaucoma onset risk
Patent: FP 1388590-A 19 11-FEB-2004;
Sysmex Corporation (JP)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.7%; Score 16.8; DB 1; Length 20; Best Local Similarity 90.0%; Pred. No. 1.1e+03; Matches 18; Conservative 0; Mismatches 2; Indels
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Reichwald, K.
Mchri variant associated with human obesity
Patent: WO 2003104489-A 82 18-DEC-2003;
Philipps-Universitaet Marburg (DE)
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Sequence 3 from Patent WO2004005541.
CO766647
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/organism="synthetic constr
/mol_type="unassigned DNA"
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Sequence 19 from Patent EP1388590.
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PAT 20-JUN-2002

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Unclassified.

1 (bases 1 to 20)

Moniab.P. and Wyatt,J.
Antisense inhibition of Phosphorylase kinase beta expression
Patent: US 6366856-A 76 09-APR-2002;
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Zhang, H. and Wyatt, J.
Antisense modulation of BH3 interacting domain death agonist
                                                                                                                                                                                                                                                                                                      Length 20;
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1 (bases 1 to 20)
Bennett, C.F., Freier, S.M. and Watt, A.T.
Antisense modulation of phospholipase A2, group VI (Ca2+-independent) expression
Patent: US 6410125-A 17 25-UNN-2002;
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Best Local Similarity 90.0%; Pred. No. 1.1e+03;
Matches 18; Conservative 0; Mismatches 2;
                                   DNA
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Patent: US 6503754-A 24 07-JAN-2003;
Location/Qualifiers
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Sequence 76 from patent US 6368856.
AR205392
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Sequence 24 from patent US 6503754.
AR271780 GI:29703348
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/organism="unknown"
/mol_type="unassigned DNA"
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Sequence 17 from patent US 6410325.
AR215876.1 GI:23314132
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/organism="unknown"
/mol_type="genomic DNA"
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//mol type="unassigned DNA"
/db xref="taxon:32500"
/note="bescription of Artificial Sequence: an artificially synthesized primer se q uence"
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                                                                                                                                                                                                                                                                                                                             Koga,H.
Primers for synthesizing full length cDNA clones and their use Patent: EP 1396543-A 4242 10-MAR-2004;
Research Association for Biotechnology (JP)
Location/Qualifiers
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/mol_type="unassigned DNA"
/db_xref="taxon:32630"
/noTe="primer for amplification of D381611"
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18; Conservative 0; Mismatches 2;
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Best Local Similarity 90.0%; Pred. No. 1.1e+03;
Matches 18; Conservative 0; Mismatches 2;
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Diagnostic test
Patent: WO 2004018711-A 17 04-MAR-2004;
University College London (GB)
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Best Local Similarity
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CQ784102
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CQ786093/c
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PAT 10-APR-2003

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Gaps

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PAT 25-SEP-2002

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Length 20;

PAT 06-AUG-2001

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DNA

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Gaps

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Length 20;

PAT 28-AUG-2001

linear

DNA

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Starling, G.C. and Pinger, J.

Novel immunoglobulin superfamily members apex-1, apex-2 and apex-3
and uses thereof
Patent: WO 0146260-A 16 28-JUN-2001;
Bristol-Myers Squibb Co. (US)
Location/Qualifiers
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sequlatory sequence for the specific expression in dendritic cells and uses thereof
Patent: WO 0151631-A 56 19-JUL-2001;
                                                                             Unknown.
Unclassified.

I (bases 1 to 20)

Monia, B.P. and Cowsert, L.M.
Antisense inhibition of telomeric repeat binding factor 2 expression
Patent: US 6300132-A 72 09-OCT-2001;
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    Sequence 72 from patent US 6300132.
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Sequence 16 from Patent W00146260.
AXI80379
AXI80379.1 GI:15132316
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Sequence 56 from Patent WO0151631.
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/organism="unknown"
/mol_type="genomic DNA"
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Monia,B.P. and Cowsert,L.M.
Antisense inhibition of telomeric repeat binding factor 2
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Patent: US 6300132-A 70 09-OCT-2001;
Location/Qualifiers
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Sequence 71 from patent US 6566135.
AR337146.1 GI:33723000
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/wol_type="genomic DNA"
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Reske-Kunz, Angelika (DE) ; Ross, Xiaolan (DE) ; Ross, Ralf (DE) Bros, Matthias (DE) Location/Qualifiers
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Detection of epigenetic abnormalities and diagnostic method basedthereon
Patent: WO 03104487-A 2 18-DEC-2003;
Centre For Addiction and Mental Health (CA)
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"Organism="synthetic construct"

/mol_type="unassigned DNA"

/db_xref="caxon:32630"

/note="primer 'Alu For' (see Example 1)"
                                                                                                                                                                                                                                                                                                         linear
                                            1. .20
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/note="artificial sequence"
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/mol_type="unassigned DNA"
/mol_txref="taxon:32630"
/noce="amorce PCR"
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Sequence 31 from Patent WO02100896.
AX657318
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Sequence 2 from Patent W003104487.
AX962284
AX962284.1 GI:40881559
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OS Unidentified

OS Unidentified

PD 2001513996-A/25

PD 11-SEP-2001

PF 18-AUG-1999 UP 2000507794

PR 22-AUG-1997 US 08/916384,06-MAY-1998 US 09/073567 PI
JTANDONG CHEN, SUDHIR AGRAWAL,RUIWEN ZHANG

PC C12N15/09, A61K31/47, A61K31/7088, A61K48/00, A61P35/00, C07H21/00,

PC C12N15/00

CC Strandedness: Both;

CC Topology: Linear;

CC Antisense oligonucleotide specific to MDM2

FF Key

Location/Qualifiers

Location/Qualifiers

FT source

/ organism='Unidentified'.
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Patent: JP 2001321190-A 1261 20-NOV-2001;
THE INSTITUTE OF PHYSICAL AND CHEMICAL RESEARCH, YUGENKAISHA
GENOTECHS
OS Artificial Sequence
PN JP 2001321190-A/1261
PD 20-NOV-2001
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0.7%; Score 16.8; DB 1; Length 20;
Best Local Similarity 90.0%; Pred. No. 1.1e+03;
Matches 18; Conservative 0; Mismatches 2; Indels
Query Match 0.7%; Score 16.8; DB 1; Length 20; Best Local Similarity 90.0%; Pred. No. 1.1e+03; Matches 18; Conservative 0; Mismatches 2; Indels
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JP 2001513996-A/25.
unidentified
unidentified
unidentified.
1 (bases 1 to 20)
Chen,J., Agrawal,S. and Zhang,R.
Antisense oligonucleotide specific to MDW2
Patent: JP 2001513996-A 25 11-SEP-2001;
                                                                                                                                                                                                                                                      BD073986 20 bp DNA 1
Antisense oligonucleotide specific to MDM2.
BD073986
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A method of arraying genome clone.
BD089017 GI:22634627
JP 2001321190-A/1261.

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synthetic construct
artificial sequences.
1 (bases 1 to 20)
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Wed Jan 26 12:47:31 2005

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DDIA 110-BD DNA 110-BT 18-SEP-2002

CON Antisense modulation of human MDM2 expression.

BD138100.1 G1:23233045

JP 2002508944-A/26.

Unidentified

unclassified.

I (Dases 1 to 20)

Miraglia,L.J., Nero,P., Graham,M.J., Monia,B.P. and Cowsert,L.M.

Antisense modulation of human MDM2 expression

LSIS PHARMACEUTICALS INC

OS Unidentified

Patent: JP 2002508944-A/26

PN JP 2002508944-A/26

PN
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C12Q1/68,
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Antisense modulation of human MDM2 expression FH
Location/Qualifiers
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JP 2002505888-A/135.
Synthetic construct
Synthetic construct
Synthetic construct
artificial sequences.
I (bases 1 to 20)
Sidlanki, D.
Detection of neoplasia by analysis of saliva
Lotenti, D.
Detection of neoplasia by analysis of saliva
Detection of neoplasia by analysis of saliva
Lotenti, D.
Detection of neoplasia by analysis of saliva
Lotenti, D.
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Detection of neoplasia by analysis of saliva.
BD134311
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90.0%; Pred. No. 1.1e+03;
tive 0; Mismatches 2;

    .20
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/db_xref="taxon:32630"

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Strandedness: Single;
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BD134311/c
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20 bp DNA linear PAT 18-SEP-2002
Primer for synthesizing full-length cDNA and use thereof.
BD128026
BD128026.1 GI:23222971
JP 2002017375-A/3457.
12-MAR-2001 JP 2001068285
EIICHI SOEDA
CI2N15/09,CI2N15/09,CI2M1/00,CI2Q1/68,G01N33/53,G01N33/566, PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Koga, H.
Primer for synthesizing full-length cDNA and use thereof
Patent: JP 2002017375-A 3457 22-JAN-2002;
Patent: JP 2002017375-A/3457
OS Unidentified
PN JP 2002017375-A/3457
PD 22-JAN-2002
PF 07-JUL-2000 JP 2000253172
PF 07-JUL-2000 JP 2000253172
PI TOSHIO OTA, TETSUO NISHIKAWA, TAKAO ISOGAI, KOJI HAYASHI, SHIZUKO
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Ota,T., Nishikawa,T., Isogai,T., Hayashi,K., Ishii,S., Kawai,Y.,
Wakamatsu,A., Sugiyama,T., Nagai,K., Kojima,S., Otsuki,T. and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PI ISHII,
PI YURI KAWAI,AI WAKAMATSU,TOMOYASU SUGIYAWA,KEIICHI NAGAI, PI
SHINICHI KOJIMA,
                                                                                                                                                       Key
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Description of Artificial Sequence:Synthetic DNA FH
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/organism='Artificial Sequence'
Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.7%; Score 16.8; DB 1;
llarity 90.0%; Pred. No. 1.1e+03;
Conservative 0; Mismatches 2;
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/db_xref="taxon:32644"

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Schumm, J.W. and Bacher, J.W.
Materials and methods for indentifying and analyzing intermediate materials and materials A standam repeat DNA markers
Patent: US 6238863-A 67 29-MAY-2001,
Location/Qualifiers
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Schumm, J.W. and Bacher, J.W.
Schumm, J.W. and Bacher, J.W.
Materials and methods for indentifying and analyzing intermediate tandem repeat DNA markers
Patent: US 6238863-A 112 29-MAY-2001;
Location/Qualifiers
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             /organism='Unidentified'.
Location/Qualifiers
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AR154062
AR154062.1 GI:15122115
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Sequence 67 from patent US 6238863.
AR154017 GI:15122070
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    .20
    /organism="unidentified"
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    .21
    organism="unknown"

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BD056594.1 GI:22602200
JP 2001508291-A/51.
synthetic construct
artificial sequences.
I (bases I to 21)
Lifton, R.P. and Simon, D.B.
Method to diagnose and treat pathological conditions resulting from deficient ion transport
Patent: JP 2001508291-A 51 26-JUN-2001;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Method to diagnose and treat pathological conditions resulting from BD056594
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         PAT 06-AUG-2001
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4P 2001508291-A/51
26-JUN-2001
19-DEC-1997 JP 1998530123
31-DEC-1997 US 08/778052
RICHARD P LIFTON DAVID B SIMON
C12N15/09, C07K14/435, C07K16/00, C12N1/15, C12N1/19, C12N1/21, PC
                                                                                                                                                                         Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                                                                                                                                                                                           Daly,M., Hudson,T.J., Lander,E.S., Rioux,J. and Siminovitch,K. Ibd.related polymorphisms
Patent: WO 1012511-A 1453 14-JUN-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH (US); Ellipsis Biotherapeutics Corporation (CA)
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            linear
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            DNA
21 bp DN Sequence 1453 from Patent WO0142511. AX183700
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/db_xref="taxon:9606"
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/organism="Homo sapiens"
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Homo sapiens
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Gaps ö PAT 29-SEP-1999

linear

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Gaps

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PAT 07-SEP-2000

linear

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Unclassified.

1 (bases 1 to 22)

1 (bases 1 to 22)

Warnsy,G.A. and Smith,M.W.

Method for generation of sequence sampled maps of complex genomes

Patent: US 5851760-A 287, 22-DEC-1998;
 JAMES W SCHUMM, JEFFREY W BACHER
C12N15/09,C12Q1/68,C12N15/00
Strandedness: Single;
Topology: Linear;
Material and method for specifying and analyzing medium-size
tandem repeat
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Unclassified.
1 (bases 1 to 22)
Chenchik,A., Jokhadze,G. and Bibilashvilli,R.
Methods of assaying differential expression
Patent: US 5994076-A 25 30-NOV-1999;
Location/Qualifiers

    .21
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                                                                                                       Location/Qualifiers
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Sequence 287 from patent US 5851760.
AR066939 GI:5998161
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Sequence 25 from patent US 5994076.
AR089905
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/mol_type="unassigned DNA"
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AR089905/c
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Patent: JP 2002502606-A 67 29-JAN-2002;
PROMEGA CORP

SO Unidentified

PN JP 2002502606-A/67

PP 39-JAN-2002

PF 04-FEB-1999 JP 2000530608

PR 04-FEB-1999 US 09/018584

PI JAMES W SCHUMM, JEFFREY W BACHER

PC C12N15/09, C12Q1/68, C12N15/00

CC Strandedness: Single;

CC Topology: Linear;

CC Material and method for specifying and analyzing medium-size
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Material and method for specifying and analyzing medium-size tandem repeat DNA marker.
BD130168 BD130168.1 GI:23225113
JP 2002502606-A/112.
                                                                                                                                        BD130123.1 GI:23225068
JP 2002502606-A/67.
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1 (bases 1 to 21)
Schumm, J.W. and Bacher, J.W.
Material and method for specifying and analyzing medium-size tandem
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Schumm, J.W. and Bacher, J.W.
Material and method for specifying and analyzing medium-size tandem
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Location/Qualifiers
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Patent: JP 2002502606-A 112 29-JAN-2002;
PROMEGA CORP

    .21
    /organism="unidentified"
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    /db_xref="taxon:32644"

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JP 2002502606-A/112
29-JAN-2002
04-FEB-1999 JP 2000530608
04-FEB-1998 US 09/01858
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PAT 15-MAY-2004

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Homo sapiens
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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1 (bases 1 to 22)

Moore, K. and Nagle, D. L.
Methods amposition for the diagnosis and treatment of body weight disorders, including obesity
Patent: US 6713277-A 93 30-MAR-2004;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Daly,M., Hudson,T.J., Lander,E.S., Rioux,J. and Siminovitch,K. Ibd-related polymorphisms
Patent: WO 0142511-A 2001 14-JUN-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH (US) ; Ellipsis
                                                                                                                                                                                              0.7%; Score 16.8; DB 1; Length 22; llarity 90.0%; Pred. No. 1.1e+03; Conservative 0; Mismatches 2; Indels
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                                     1 (bases 1 to 22)
Chenchik,A., Jokhadze,G. and Bibilashvilli,R.
Methods of assaying differential expression
Patent: US 648945-A 25 03-DEC-2002;
Location/Qualifiers
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Best Local Similarity 90.0%; Pred. No. 1.1e+03;
Matches 18; Conservative 0; Mismatches 2;
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Sequence 2001 from Patent WO0142511.
AX184248
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Sequence 93 from patent US 6713277.
AR490692
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/mol_type="unassigned DNA"
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                                                                                                                                          /organism="unknown"
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/organism="unknown"
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                     Unclassified.
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Moore, K. and Nagle, D.Lynn.
Methods and compositions for the diagnosis and treatment of body
weight disorders, including obesity
Patent: US 6274339- A 92 14-AUG-2001;
Location/Qualifiers
                       Gaps
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3 1 (bases 1 to 22)
5 Chenchik, A., Jokhadze, G. and Bibilashvilli, R. Methods of assaying differential expression
AL Patent: US 632829-A 25 05-MAR-2002;
Location/Qualifiers
 90.0%; Pred. No. 1.1e+03; ive 0; Mismatches 2;
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    .22
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/mol_type="unassigned DNA"

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/organism="unknown"
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AR164891
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Sequence 25 from patent US 6352829.
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Best Local Similarity 90.0
Matches 18; Conservative
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THE INSTITUTE OF PHYSICAL AND CHEMICAL RESEARCH, YUGENKAISHA
GENOTECHS
OS Artificial Sequence
PN 72001321190-A/1521
PD 2001321190-A/1521
PP 12-MAR-2001 JP 2001068285
PF 12-MAR-2001 JP 2001068285
PF 18-IICHI SOEDA
PC C12N15/09,C12N15/09,C12M1/00,C12Q1/68,G01N33/53,G01N33/5
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Pred. No. 1.1e+03;
0; Mismatches 1; Indels
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85.7%; Pred. No. 1.1e+03; indels ive 0; Mismatches 3; Indels
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Monia B.P., Baker B.F. and Cowsert, L.M.
Antisense modulation of CREL expression
Patent: US 601652-A 45 14-DEC-1999;
Location/Qualifiers
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AR094543
AR094543.1 GI:10021565
                                                                                                                         A method of arraying genome clone. BD089277

    .22
    /organism="synthetic col/mol_type="genomic DNA"
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                                     2089 TTATTTTTGAGACCGAGT 2109
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JP 2001321190-A/1521.
synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 22)
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ilarity 90.0%;
Conservative 0
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Best Local Similarity 94.4
Matches 17; Conservative
Best Local Similarity 85.7
Matches 18; Conservative
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synthetic construct
artificial sequences.

1 (bases 1 to 18)
Shiozawa, S., Komai, K., Yaqi, H. and Matsuura, N.
Shiozawa, S., Komai, K., Yaqi, H. and Matsuura, N.
Genomic DNA participating in rheumatoid arthritis, method of diagnosing the same, method of judging the onset risk of the same and diagnostic for detecting the same patent: MO 02019466-A 10 10-OCT-2002;
SHUNICHI SHIOZAWA, KOICHIRO KOMAI, HIROFUMI YAGI, NAO MATSUURA
OS Artificial Sequence
PN WO 02019466-A/10
PD 10-OCT-2002
PP 29-MAR-2002 WO 2002JP003191
PR 30-MAR-2001 JP DIP 102006
PI SHUNICHI SHIOZAWA, KOICHIRO KOMAI, HIROFUMI YAGI, NAO MATSUURA PC
C12N15/09,C12Q1/68,G01N33/56,G01N33/50
CC Synthesized oligomucleotide
                                                                                                                  PAT 16-APR-2003 method of
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                                                                             Genomic DNA participating in rheumatoid arthritis, method of diagnosing the same, method of judging the onset risk of the same and diagnostic for detecting the same.
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0; Mismatches 1;
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/mol_type="unassigned DNA"
/db xref="taxon:32630"
/noTe="Priner"
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Sequence 993 from Patent WO0129262.
AX115870.1 GI:14032812
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WO 02079466-A/10.
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PAT 11-MAY-2001

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SOURCE

FEATURES

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Homo sapiens
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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0.7%; Score 16.4; DB 1;
Best Local Similarity 94.4%; Pred. No. 1.1e+03;
Matches 17; Conservative 0; Mismatches 1;
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/nofe="primer"
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Seguence 3594 from Patent WO0129262.
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Sequence 3358 from Patent WO0129262.
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Sequence 8 from Patent WO0222879.
AX412182
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 0.7%; Score 16.4; DB 1; Length 18; 94.4%; Pred. No. 1.1e+03; ive 0; Mismatches 1; Indels
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/mol_type="unassigned DNA"
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Sequence 1786 from Patent WO0129262.
AX116663.1 GI:14033605
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                                                                       2174 CCGGGTTCGCACCATTCT 2191
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Bacher, J.W., Flanagan, L. and Nassif, N.

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Detection of microsatellite instability and its use in diagnosis of
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Parent: WO 02077272-A 82 03-OCT-2002;
Epigenomics AG (DE)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Vanilloid receptor-related nucleic acids and polypeptides
Patent: WO 02101045-A 46 19-DEC-2002;
Novartis AG (CH) ; IRM LLC (BM)
Location/Qualifiers
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14.4%; Pred. No. 1.1e+03;
ve 0; Mismatches 1; Indels
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Sequence 82 from Patent W002077272.
AX598742
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/note="MONO-15 primer"
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Patent: WO 0222879-A 8 21-MAR-2002;
PROMEGA CORPORATION (US)
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/organism="Homo sapiens"

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                                                                                                                                                                                                                                             2152 CTCACTGCAAGCTCTGCC 2169
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artificial sequences.
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Best Local Similarity 94.4%;
Matches 17; Conservative
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Homo sapiens
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Best Local Similarity 94.4*
Matches 17; Conservative
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AX708864/c
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/mol type="genomic DNA"
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/note="Description of Combined DNA/RNA Molecule:Synthetic Oligomer Sequence-Synthetic Probe Sequence"
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/db xref="taxon:325830"
/note="Description of Combined DNA/RNA Molecule:Synthetic Oligomer Sequence-Synthetic Probe Sequence"
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Methods, kits and compositions pertaining to the suppression of
detectable probe binding to randomly distributed repeat sequences
in genomic nucleic acid
Patent: WO 03027328-A 16 03-APR-2003,
Boston Probes, Inc. (US); DakoCytomation Denmark A/S (DK)
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0.7%; Score 16.4; DB 1; Length 18;
Best Local Similarity 94.4%; Pred. No. 1.1e+03;
Matches 17; Conservative 0; Mismatches 1; Indels
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Pred. No. 1.1e+03;
0; Mismatches 1;
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/db_xref="taxon:32630"
/note="Oligonucleotide primer"
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Sequence 4 from Patent WO03027328.
AX741030
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Best Local Similarity 94.4%;
Matches 17; Conservative
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artificial sequences.
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Homo sapiens (human)
Homo sapiens
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FH Key
FT source
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BD102660/c
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MATERIALS AND METHODS RELATING TO THE IDENTIFICATION AND SEQUENCING
OF THE BRCA2 CANCER SUSCEPTIBILITY GENE AND USES THEREOF
PATENT: WO 9719110-A 124 29-MAY-1997;
CANCER RES CAMPAIGN TECH (GB)
Other publication AN 7635096 19970611
Other publication GB 2307477 19970528.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Patent: WO 0142511-A 1454 14-JUN-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH (US) ; Ellipsis
Biotherapeuties Corporation (CA)
Location/Qualifiers
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 Score 16.4; DB 1; Length 18;
Pred. No. 1.1e+03;
0; Mismatches 1; Indels
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Sequence 124 from Patent.W09719110.
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Query Match
Best Local Similarity 94.4%;
Matches 17; Conservative (
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          PAT 06-AUG-2001
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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synthetic construct
artificial sequences.

E 1 (bases 1 to 19)
S Chimera animal
L Patent: WO 0187059-A 2 22-NOV-2001;
JAPAN SCIENCE AND TECHNOLOGY CORP, CHISE MUKAIDANI, KATSUTOSHI
YOSHIZATO, TOSHINORI FURUKAWA
OS Artificial Sequence
PN WO 0187059-A/2
PN WO 0187059-A/2
PN WO 10187059-A/2
PN RAY-2001 WO 2001JF004193
PR 19-MAY-2000 JP 00P 149079
PR 19-MAY-2000 JP 00P 149079
PR 19-MAY-2000 JP 00D 149079
PR 19-MAY-2000 JP 00D ATTIFICIAL SEQUENCE: Synthesized CC
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Patent: WO 0142511-A 1653 14-JUN-2001;
WHITEHEAD INSTITUTE FOR BIOMEDICAL RESEARCH (US); Ellipsis Blotherapeutics Corporation (CA)
Location/Qualifiers
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0.7%; Score 16.4; DB 1; Length 19;
Best Local Similarity 94.4%; Pred. No. 1.1e+03;
Matches 17; Conservative 0; Mismatches 1; Indels
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    .19
/organism='Artificial Sequence'

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/organism="synthetic construct"
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            DNA
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19 bp DN Sequence 1653 from Patent W00142511. AX183900

    19 /organism="Homo sapiens"
/mol_type="unassigned DNA"
/db_xref="taxon:9606"

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/db_xref="taxon:32630"
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PAT 08-SEP-2000

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Gaps

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PAT 16-MAY-2001

DEFINITION ACCESSION VERSION KEYWORDS SOURCE ORGANISM

AUTHORS TITLE JOURNAL

COMMENT

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RESULT 1044

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BD137510/c LOCUS

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Unclassified.

1 (bases 1 to 20)
Lalouel,J.-M., Jeunemaitre,X., Lifton,R.P., Soubrier,F.,
Kotelevtsev,Y. and Corvol,P.
Method to determine predisposition to hypertension
Patent: US 5998145_A 19 07-DEC-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 (bases 1 to 20)
Lalouel,J.-M., Jeunemaitre,X., Lifton,R.P., Soubrier,F.,
Kotelevtsev,Y. and Corvol,P.
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1 (bases 1 to 20)

Lalouel,J.-M. and Jeunemaitre,X.

Method to determine predisposition to hypertension
Patent: US 6153386-A 19 28-NOV-2000;
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0.7%; Score 16.4; DB 1;
Best Local Similarity 94.4%; Pred. No. 1.1e+03;
Matches 17; Conservative 0; Mismatches 1;
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Sequence 19 from patent US 6165727.
AR122443.
AR122443.1 GI:14106760
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Sequence 19 from patent US 6153386.
AR119526
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/organism="unknown"
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/wol_type="unassigned DNA"
                                                                              20 bp | Sequence 19 from patent US 5998145.
AR092309
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 18 CCGAGGCTGGAGTGCAGT 1
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                                                                               PAT 18-SEP-2002
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BD137510.1
BD137510.1
BD137510.1
GT:23232455
JP 2002045087-A/2.
Synthetic construct
synthetic construct
artificial sequences.
1 (bases 1 to 19)
Mukaidani, C., Yoshizato, K. and Furukawa, T.
Chimera animal
Patent: JP 202045087-A 2 12-FEB-2002;
PAPAN SCIENCE AND TECHNOLOGY CORP, HIROSHIMA INDUSTRIAL TECHNOLOGY
                                                                                                                                                                                                                                                                                                                                                            PF 18-MAY-2001 JP 2001150098
PI CHISE MUKAIDANI,KATSUTOSHI YOSHIZATO,TOSHINORI FURUKAWA PC
A01K67/027,C12N15/09,C12Q1/02,G01N33/15,G01N33/50//(C12Q1/02, PC
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1 (bases 1 to 20)
Lalouel,J.M., Jeunemaitre,X., Lifton,R.P., Soubrier,F., Koteleytsev, Y. and Corvol,P.
Method to determine predisposition to hypertension
Patent: US 5763168-A 19 09-JUN-1998;
Location/Qualifiers
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PC C12N15/00

C12N15/00

Oligonucleotide Location/Qualifiers

FT Source | 1.19
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Pred. No. 1.1e+03;
0; Mismatches 1; Indels
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/organism="synthetic construct"
/mol_type="genomic DNA"
/db_xref="taxon:32630"
                                                                                 DNA
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Sequence 19 from patent US 5763168.
AR011709
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/mol_type="unassigned DNA"
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JP 2002045087-A/2
12-FEB-2002
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Best Local Similarity 94.4%;
Matches 17; Conservative
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Best Local Similarity 94.4
Matches 17; Conservative
                                                                                               Chimera animal.
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FEATURES

RESULT 1045

8 g AR011709/c LOCUS

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ACCESSION VERSION KEYWORDS SOURCE

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Gaps

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PAT 16-MAY-2001

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TITLE JOURNAL FEATURES

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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.

E. 1 (Abases 1 to 20)

E. 1 (Abases 1 to 20)

Method of quantifying hypertensive constitution

L. Patent: JP 200219012-A 19 02-ULL-2002;
UNIVERSITY OF UTAH RESEARCH FOUNDATION

OS Homo sapiens (human)

PN JP 2002519012-A/19

PP 15-APR-1999 JP 2000557000

PR 29-UTM-1998 US 09/106216

PI JEAN MARC LALOUEL, XAVIER JEUNEMAITRE

PC C12Q1/68, C12N15/99, C12N15/90

CC Method of quantifying hypertensive constitution FH Key
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JEAN MARC LALOUEL, XAVIER JEUNEMAITRE
C1201/68, C12N15/09, C12N15/00
Method of quantifying hypertensive constitution FH
Location/Qualifiers
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/organism='Homo sapiens (human)'.
Location/Qualifiers
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Pred. No. 1.1e+03;
0; Mismatches 1; Indels
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14.4%; Pred. No. 1.1e+03;
ve 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                       BD217343 20 bp DNA linear Method of quantifying hypertensive constitution.
Method to determine predisposition to hypertension Patent: US 6165727-A 19 26-DEC-2000; Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Unidentified
unidentified
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unclassified.
1 (bases 1 to 20)
Hirotsu, T., Karsahi, H., Matsuhisa, A. and Ono, N.
PROBE FOR DIAGNOSIS OF INFECTIOUS DISEASE
PATH: JP 1994133798-A 5 17-MAY-1994;
FUSO YAKUHIN KOGYO KK, ONO NORIYA
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    .20
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    /mol_type="unassigned DNA"

    .20
    /organism="Homo sapiens"
    /mol_type="genomic DNA"
    /db_xref="taxon:9606"

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JP 2002519012-A/19.
Homo sapiens (human)
Homo sapiens
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E07490
E07490.1 GI:2175628
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Best Local Similarity 94.4%;
Matches 17; Conservative
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Best Local Similarity 94.4%;
Matches 17; Conservative
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Artificial sequences.

N JP 1994133798-A/5

N JP 1994133798-A/5

N JP 1994133798-A/5

F 23-OCT-1992 JP 1992285802

I HIROTSU TAKUO, KARASHI HIROYUKI, MATSUHISA AKIO, ONO NORIYA PC C12Q1/68, C12Q1/04, (C12Q1/04, C12R1:725);

E strandedness: Single;

C topology: Linear;

C hypothetical: No;
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Lalouel, J.-M., Jeunemaitre, X., Lifton, R.P., Soubrier, F.,
Kotelevtsav, Y. and Corvol, P.
Angiotensinogen gene variants and predisposition to hypertension
Patent: US 5589584-A 19 31-DEC-1996;
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/organism='Artificial sequences' FT
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Pred. No. 1.1e+03;
0; Mismatches 1; Indels
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Unclassified.
(Dases 1 to 20)
Ohno,T., Hirotsu,T., Keshi,H. and Matsuhisa,A.
                                                                                                                                                                                                                                                                                   /note='Probe for PCR
                                                                                                                                                                                          Location/Qualifiers
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Sequence 19 from patent US 5589584.
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Sequence 5 from patent US 5708159.
I80108.1 GI:3208398
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/organism="unidentified"
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/organism="unknown"
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Best Local Similarity 94.4%;
Matches 17; Conservative
                                                                                                                                                                      anti-sense: No;
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                                                                                                            CC stranded
CC topology:
CC anti-sens
FH Key
FT source
FT source
FT
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Unclassified.
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PAT 12-JAN-2001

Bource

JOURNAL

TITLE

FEATURES

RESULT 1053 AR492707/c

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DEFINITION

ACCESSION

VERSION

KEYWORDS

SOURCE

ORGANISM

REFERENCE AUTHORS JOURNAL

TITLE

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synthetic construct
artificial sequences.

E 1 (bases 1 to 21)
SYNTHetic construct
artificial sequences.

E 1 (bases 1 to 21)
SY Yoshikawa, T. and Hattori, B.

Polymorphism of upstream region of human cholecystokinin gene, identification method and reagent thereof, and method for diagnosis of anxiety discorders based thereon
of anxiety discorders based thereon
of anxiety discorders based thereon
Datent: JP 2002171990-A 5 18-JUN-2002;
THE INSTITUTE OF PHYSICAL AND CHEMICAL RESEARCH
OS Artificial Sequence
PN JP 2002171990-A/5
PD 18-JUN-2002
PD 18-JUN-2002
PD 09-DEC-2000 JP 2000375090
PI TAKES VOSHIKAWA, EIJI HATTORI
PC C12N15/09, C12Q1/68, G01N33/556, C12N15/09 CC
Description of Artificial Sequence: upstream primer p5 FH Key
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             21 bp DNA linear PAT 17-JAN-2003 Polymorphism of upstream region of human cholecystokinin gene, identification method and reagent thereof, and method for diagnosis BDIG anxiety disorders based thereon.
                                                                                                                                                                                                                                                                                            Shimkets,R.A., Fernandes,E. and Boldog,F.
Secreted polypeptides and corresponding polynucleotides
Patent: WO 0070046-A 47 23-NOV-2000;
Curagen Corporation (US)
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/organism='Artificial Sequence'.
Location/Qualifiers
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O.7%; Score 16.4; DB 1;
Best Local Similarity 94.4%; Pred. No. 1.1e+03;
Matches 17; Conservative 0; Mismatches 1;
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0.7%; Score 16.4; DB 1;
Best Local Similarity 94.4%; Pred. No. 1.1e+03;
Matches 17; Conservative 0; Mismatches 1;
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                                                                                                             DNA
                                                                                                         AX050293 21 bp
Sequence 47 from Patent WO0070046.
AX050293
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        1 ATTACAGGCGTGAGCCAC 18
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JP 2002171990-A/5.
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Probe for diagnosing infectious diseases which hybridizes with DNA from candida albicans
Patent: US 5708159-A 5 13-JAN-1998;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.7%; Score 16.4; DB 1; Length 20; Best Local Similarity 94.4%; Pred. No. 1.1e+03; Matches 17; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                     Length 20;
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94.4%; Pred. No. 1.1e+03;
rative 0; Mismatches 1;

    .20
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    /note="Primer"

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Patent: US 6716975-A 77 06-APR-2004;
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Sequence 2886 from Patent W00129262.
AX117763
                                                                                                     /organism="unknown"
/mol_type="unassigned DNA"
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Sequence 77 from patent US 6716975.
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/organism="unknown"
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                                                                Location/Qualifiers
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                                                                                                                                                                                                                                                   393 GTTAGACCAAAGCCATTG 410
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Best Local Similarity 94.4%
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Best Local Similarity 94.4
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Unknown.
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source

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RESULT 1054

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AX117763

DEFINITION

ACCESSION VERSION KEYWORDS SOURCE

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RESULT 1057

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ACCESSION VERSION KEYWORDS SOURCE

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rITON Multiple amplification of short tandem repeat gene site.

Multiple amplification of short tandem repeat gene site

Multiple amplification of short tandem repeat gene site
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Homo sapiens (human)
DP 2002530121-A/21
17-SEP-2002
24-NOV-1999 UP 2000584113
25-NOV-1999 US 09/199542
JAMES W SCHUMM, CYNTHIA J SPRECHER
C1201/68, C12N15/09, C12N15/09, G01N33/58, C12N15/00,
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/organism='Homo sapiens (human)'.
Location/Qualifiers
                                                                                                                                                                         Query Match

0.7%; Score 16.2; DB 1; Length 21;
Best Local Similarity 85.7%; Pred. No. 1.2e+03;
Matches 18; Conservative 0; Mismatches 3; Indels
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 1 (bases 1 to 21)
Schumm, J.W., Micka, K.A. and Rabbach, D.R.
Multiplex amplification of short tandem repeat loci
Patent: US 5843660-A 21 01-DEC-1998;
Location/Qualifiers
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Sequence 25 from Patent WO2004003550.
CQ760468 CQ760468. GI:44903981
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Cell lines from a transgenic mouse which express biologically
active IX factor
Patent: US 5814716-A 6 29-SEP-1998;
Location/Qualifiers
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Location/Qualifiers
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Sequence 21 from patent US 5843660.
AR061829
AR061829.1 GI:5989520
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RESULT 1058

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Matches

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PAT 03-MAR-2004

RESULT 1059 AR061829/c

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artificial sequences.
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Disease risk estimating method using sequence polymorphisms in a
specific region of chromosome 19
Patent: WO 204001229-A 9 08-JAN-2004;
Aarhus University (DK); Arbejdsmilj Instituttet (National
Institute of Occupational Health) (DK)
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Disease risk estimating method using sequence polymorphisms in a
specific region of chromosome 19
Patent: WO 2004001329-A 135 08-JAN-2004;
Aarhus University (DK); Arbejdsmilj Instituttet (National
Institute of Occupational Health) (DK)
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Leyland-Jones, B.
Individualization of therapy with anticoagulants
Patent: Wo 2004003550-A 25 08-JAN-2004;
Xanthus Life Sciences, Inc. (US)
Location/Qualifiers
                                                                                         /organism="synthetic construct"
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                                                                                                                                                                      Score 16.2; DB 1;
Pred. No. 1.2e+03;
0; Mismatches 3;
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/mol_type="unassigned DNA"
/mb_xref="taxon:32630"
/not=="probe"

    .21
    /organism="synthetic construct"

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CQ760693
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Matches 18; Conserv
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DAI ICHI SBIYAKU CO LTD
OS Artificial genee

C Artificial genee

C Artificial sequence; Genes.

C Artificial sequence; Genes.

S Rattus SP. (rat)

PD 1992148678-A/1

PD 21-MAY-1992

PP 12-OCT-1990 JP 1990274194

PI SAKANO KATSUICHI, FUJIWARA HIROYUKI, AZUMABASHI NOBUYUKI, PI
MARUMOTO YAGUMASA, SATO YOSHIO

PC C12N1/21,CO7K7/10,C12N15/18//A61K37/02,C12P21/02, (C12N1/21, PC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Nucleic acid amplification primers for pcr-based clonality studies Patent: WO 2004033728-A 114 22-APR-2004;
Erasmus Universiteit Rotterdam (NL); Van Dongen, Jacobus, Johannes,
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(+1224)"
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1 (bases 1 to 21)
Sakano,K., Fujiwara,H., Azumabashi,N., Marumoto,Y. and Sato,Y.
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Synthetic DNA sequence of rat IL-1 alpha PCR primer.
E01635
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0.7%; Score 16.2; DB 1;
Best Local Similarity 85.7%; Pred. No. 1.2e+03;
Matches 18; Conservative 0; Mismatches 3;
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/mol_type="unassigned DNA"
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2270 GACAGGGTTTCACCGTGTTAG 2290
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  /note="Primer"
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                                                                                                   _RNA 1. 21
/gene='rat IL-1 alpha PCR primer'
Location/Qualifiers
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Schumm, J.W. and Sprecher, C.J.
Multiplex amplification of short tandem repeat loci
Patent: US 6479235-A 21 12-NOV-2002;
Location/Qualifiers
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Sequence 2381 from Patent WO0129262.
AX117258
C12R1:19),
(C12P21/02,C12R1:19),C07K99:00;
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AR252820
AR252820.1 GI:27301169
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                          strandedness: Single;
topology: Linear;
hypothetical: No;
anti-sense: No;
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synthetic construct
artificial sequences.
                                                                                                         misc_RNA
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PAT 11-MAY-2001
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                  Gaps
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Query Match 0.7%; Score 16.2; DB 1; Length 21; Best Local Similarity 85.7%; Pred. No. 1.2e+03; Matches 18; Conservative 0; Mismatches 3; Indels
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Individualization of therapy with antipsychotics
Patent: WO 02073196-A 25 19-SEP-2002;
MGGLL UNIVERSITY (CA)
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Messiaen,L. and Callens,T.
Improved mutation analysis of the nfl gene
Patent: WO 0129251-A 58 26-APR-2001;
UNIVERSITEIT GENT (BE)
Location/Qualifiers
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Sequence 58 from Patent W00129251.
AX119401
AX119401.1 GI:14036320
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Sequence 25 from Patent W002073196.
AX546456
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Leyland-Jones, B.
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AX591117
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                              linear
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Individualization of therapy with antidepressants
Patent: WO 02073197-A 25 19-SEP-2002;
MCGILL UNIVERSITY (CA)
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Patent: WO 02073206-A 25 19-SEP-2002;
McGILL UNIVERSITY (CA)
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Sequence 25 from Patent WO02073205.
AX557406
                           AX557297 21 bp 1
Sequence 25 from Patent WO02073197.
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Sequence 25 from Patent W002073206.
AX557381.1 GI:25900290
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AX557297.1 GI:25900251
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RESULT 1070
AX557297
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AX557406
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Individualization of therapy with gastroesophageal reflux disease
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Metabolic phenotyping in therapy with immunosuppressants Patent: WO 02073205-A 25 19-SEP-2002; McGILL UNIVERSITY (CA)
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/b xref="taxon:32630"
/noTe="Sequence to be used as a primer"
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// Organism="synthetic construct"
// nol_type="unassigned DNA"
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// note="Sequence to be used as a Primer"
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McGILL UNIVERSITY (CA)
Location/Qualifiers
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AX592507
AX592507.1 GI:27950585
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Query Match
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AX601690
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X593485
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        0.7%; Score 16.2; DB 1; Length 21;
85.7%; Pred. No. 1.2e+03;
tive 0; Mismatches 3; Indels
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Sequence 23 from Patent WO02084288.
AX593010
AX593010.1 GI:27950854
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           Query Match
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AX593010
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Individualization of therapy with analgesics
Patent: WO 0209094-A 25 14-NOV-2002;
MCGILL UNIVERSITY (CA)
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85.7%; Pred. No. 1.2e+03;
rative 0; Mismatches 3;
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/note="primer"
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Sequence 25 from Patent WO02088714.
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Sequence 25 from Patent W002090994.
AX597480
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Sequence 25 from Patent W002093162.
AX601690 GI:28401735
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Location/Qualifiers
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AX696046
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ACCESSION
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Tchernev,V.T., Colman,S.D., Ballinger,R.A., Padigaru,M.,
Toholenc,A.R., Shenoy,S.G., Edinger,S.R., Gerlach,V., Gangolli,E.A.,
Macdougall,J.R., Smithson,G., Peyman,J.A., Stone,D.J., Gunther,E.,
Bllerman,K., Grosse,W.M., Alsobrook,J.P., Lepley,D.M. and
Burgess,C.E.
Gpcr-like protein and nucleic acids encoding same
Patent: WO 0240539-A 137 23-MAY-2002;
Curagen Corporation (US)
Location/Qualifiers
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Individualization of therapy with hyperlipidemia agents
Patent: WO 02095402-A 23 28-NOV-2002;
MCGILL UNIVERSITY (CA)
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                                                                                                                                                    Length 21;
Individualization of therapy with antibiotic agents Patent: WO 02093162-A 25 21-NOV-2002; McGILL UNIVERSITY (CA)
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Pred. No. 1.2e+03;
0; Mismatches 3;

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    /db xref="texon:32630"
    /noTe="Primer"

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AX642809
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Individualization of therapy with alzheimer's disease agents
Patent: WO 02099422-A 25 12-DEC-2002;
MGGILL UNIVERSITY (CA)
Location/Qualifiers
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                                                                                      Query Match 0.7%; Score 16.2; DB 1;
Best Local Similarity 85.7%; Pred. No. 1.2e+03;
Matches 18; Conservative 0; Mismatches 3;
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0.7%; Score 16.2; DB 1;
Best Local Similarity 85.7%; Pred. No. 1.2e+03;
Matches 18; Conservative 0; Mismatches 3;
/mol_type="unassigned DNA"
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/note="oligonucleotide primer"
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Sequence 23 from Patent WO03008637.
AX696046
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Sequence 25 from Patent WO02099422.
AX643865
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Individualization of therapy with antiviral agents
Patent: WO 03046559-A 25 05-JUN-2003;
Xanthus Life Sciences, Inc. (US)
Location/Qualifiers
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/db_xref="taxon:32630"
/noEe="Pximer"
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                               AX773023 21 bp
Sequence 25 from Patent W003046583.
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Sequence 25 from Patent WO03046559.
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AX781407.1 GI:32698408
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Taylor,K.D., Scheuner,M.T., Rotter,J.I. and Yang,H.
Genetic test to determine non-responsiveness to statin drug
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0.7%; Score 16.2; DB 1; Length 21;
Best Local Similarity 85.7%; Pred. No. 1.2e+03;
Matches 18; Conservative 0; Mismatches 3; Indels
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                          Leyland-Jones, B.
Individualization of therapy with antihistamines
Patent: WO 03052123-A 25 26-JUN-2003;
Xanthus Life Sciences, Inc. (US)
Location/Qualifiers
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/note="Sequence to be used as a Primer"
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Individualization of therapy with anxiolitics
Patent: WO 03067257-A 25 14-AUG-2003;
MCGILL UNIVERSITY (CA)
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/mol_type="unassigned DNA"
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Sequence 25 from Patent WO03067257.
AX815463
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ARI71182.1 GI:17910132
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Location/Qualifiers
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CQ798656.1 GI:46427018
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BD203034. 1 GI:33012804
JP 2002509721-A/6060.
Homo sapiens (human)
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PN JP 2002569721-A/6060
PD JP 202569721-A/6060
PF 24-MAR-1999 JP 2000541291
PR 27-MAR-1998 US 60/079678
PI PAMELA A PAVCO, ELISABETH ROBERTS, THALE JARVIS, CLAIRE COESHOTT, PI JAMES A MCSWIGGEN
PI JAMES A MCSWIGGEN
PI JAMES A MCSWIGGEN
C12N15/09, A61R31/7088, A61R31/7125, A61K48/00, A61P3/10, A61P17/06, PC
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Pavco, P.A., Roberts, E., Jarvis, T., Coeshott, C. and Mcswiggen, J.A. Method and reagent for treating diseases or conditions concerning molecule participating in vasculogenic response Patent: JP 2002509721-A 6060 02-APR-2002; RIBOZYME PHARMACEUTICALS INC
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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0.7%; Score 16; DB 1; Length 17;
100.0%; Pred. No. 1.1e+03;
ive 0; Mismatches 0; Indels
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participating in vasculogenic response
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100.0%; Pred. No. 1.1e+03;
ive 0; Mismatches 0;

    .17
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/mol_type="genomic RNA"
/db_xref="taxon:9606"

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Galvin, M., Miller, A., Penny, L. and Riedy, M.
Genotype determination of human UDP-glucuronosyl transferase 284 (UGT2815), and 2815 (UGT2815) genes
L 97 (UGT2815) and 2815 (UGT2815) genes
Patent: JP 2002521067-A 12 16-JUL-2002;
AXYS PHARMACEUTICALS INC
OS Homo sapiens (human)
PN JP 2002521067-A/12
PD 16-JUL-2002
PP 22-JUL-1999 JP 2000562588
PR 28-JUL-1999 JP 2000562588
PR 28-JUL-1998 JP 2000562588
PR 28-JUL-1998 JP 200056288
PR 28-JUL-1998 JP 2000562
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Genetic test to determine non-responsiveness to statin drug
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Cedars-Sinai Medical Center (US)
Location/Qualifiers
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Seguence 13 from patent US 6586175.
AR349392
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Sequence 91 from Patent EP1408121.
CQ798656
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PAT 31-MAR-2003

linear

DNA

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Homo sapiens
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
mdz12
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                       Query Match
0.7%; Score 16; DB 1; Length 17;
Best Local Similarity 100.0%; Pred. No. 1.1e+03;
Matches 16; Conservative 0; Mismatches 0; Indels
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Acomica, Inc. (US)
Location/Qualifiers
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Aeomica, Inc. (US)
Location/Qualifiers
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Sequence 5447 from Patent EP1281758.
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AX692715.1 GI:29415673
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/mol_type="unassigned DNA"
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                                     1 (bases 1 to 17)
Galvin,M., Miller,A., Penny,L. and Riedy,M.
Genotyping the human UDP-glucuronosyltransferase 2B7 (UGT2B7) gene
Patent: US 6586175-A 13 01-JUL-2003;
Location/Qualifiers
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Mammalla, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
mdz12
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Aeomica, Inc. (US)
Location/Qualifiers
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AX692567
AX692567.1 GI:29415525
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Cedars-Sinai Medical Center (US)
Location/Qualifiers

    .17
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/db_xref="taxon:9606"

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Sequence 91 from Patent WO0102606.
AX068540
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/organism="unknown"
/mol_type="genomic DNA"
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Four human zinc-finger-containing proteins : mdz3, mdz4, mdz7 and
mdz12
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                     Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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Aeomica, Inc. (US)
Location/Qualifiers
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Molecular Engines Laboratories (FR)
Location/Qualifiers
                                           AX692741 17 bp D)
Sequence 5473 from Patent EP1281758.
AX692741
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Seguence 986 from Patent WO03025175.
AX729352
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Sequence 34 from patent US 6171833.
AR123981
AR123981.1 GI:14109342
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    .17
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Homo sapiens
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Unclassified.
1 (bases 1 to 19)
Sinskey, A.J., Lessard, P.A. and Willis, L.B.
Pyruvate carboxylase polypeptide from Corynebacterium glutamicum
Patent: US 6403351-A 34 11-JUN-2002;
Location/Qualifiers
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[bases 1 to 19]
Sinskey, A.J., Lessard, P.A. and Willis, L.B.
Pyruvate carboxylase from corynebacterium glutamicum
Patent: 18 6171833-A 34 09-JAN-2001;
Location/Qualifiers
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Unclassified.
3 1 (bases 1 to 19)
5 0 sowski,b.
In vivo assay for intravasation
AL Patent: US 6228345-A 2 08-MAY-2001;
Location/Qualifiers
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Sequence 34 from patent US 6403351.
AR213180
                                                                                                                               /organism="unknown"
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Sequence 2 from patent US 6228345.
AR148945.1 GI:15113536
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/organism="unknown"
/wol_type="genomic DNA"
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Search completed: January 25, 2005, 16:03:02
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A novel nucleic acid molecule
Patent: WO 9851790-A 2 19-NOV-11998;
CANCILLA MICHAEL ROBERT (AU); CHOO KONG HONG ANDY (AU); SART
DESIREE DU (AU); AMRAD OPERATIONS PTY LID (AU)
Location/Qualifiers
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Unclassified.
1 (bases 1 to 20)
Monia, B.P. and Cowsert, L.M.
Antisense inhibition of telomeric repeat binding factor 2
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Best Local Similarity 100.0%; Pred. No. 1.2e+03;
Matches 16; Conservative 0; Mismatches 0;
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AX162688
AX162688.1 GI:14544019
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Patent: US 6300132-A 64 09-OCT-2001;
Location/Qualifiers
1. .20
/organism="unknown"
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    .19
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Sequence 64 from patent US 6300132.
AR370243. GI:34606749
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Sequence 2 from Patent WO9851790.
AX033910
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Homo sapiens
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Mammalia; Butheria; Primates; Catarrhini; Hominidae; Homo.
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Accession number cg44911343"
                                                                                                         Shimkets, R.A. and Leach, M.

Nucleic acids containing single nucleotide polymorphisms and methods of use thereof

Patent: WO 014021-A 6016 07-JUN-2001;

Curagen Corporation (US)

Location/Qualifiers
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Best Local Similarity 62.5%; Pred. No. 1.1e+03;
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Patent No. 6783961
Patent No. 6783961
APPLICANT: Dunmas Milne Edwards, J.B.
APPLICANT: Dunmas Milne Edwards, J.B.
APPLICANT: Dunmas Milne Edwards, J.S.
APPLICANT: Optidano, J.Y.
TITLE OP INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
Patent No. 6783961
FILE REFERENCE: 59.022.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
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Pred. No. 0.017;
0; Mismatches 5; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 13007, Application US/09621976

Patent No. 6639063

GENERAL INFORMATION:
APPLICANT: Jobert, S.
TITLE OF INVENTION:
TITLE OF INVENTION:
CURRENT APPLICANT: OLOGHER: US/09/621,976

CURRENT FILING DATE: 2000-07-21

NUMBER OF SEQ ID NOS: 19335

SOFTWARE: Patent.pm

SEQ ID NO 13007
                                                                                                                                                                                                                                                                                                                                                                                                                                                             2293 AGGATGGTCTCGATCTCCTGACCTCGTGATCCGCCC 2328
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       63 AGGATGGTCTTGATCTCCTGACCTCGTGATCCGCCC 98
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FILE REFERENCE: 59.US2.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
FRIOR APPLICATION NUMBER: US 60/122,487
FRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SQQ ID NO 18683
LENGTH: 100
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Best Local Similarity 94.69
warches 88; Conservative
                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 93.8
Matches 90; Conservative
                                                                                                                                                                                                                  CRGANISM: Homo sapiens US-09-513-999C-18683
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US-09-621-976-13007
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2263 TAGTAGAGACAGGGTTTCACCGTGTTAGCCAGGATGGTCTCGATCTCCTGACCTCGTGAT 2322
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US-09-513-999C-32786

Sequence 32786, Application US/09513999C

Patent No. 6783961

GENERAL INFORMATION:

APPLICANT: Dunas Milne Edwards, J.B.

APPLICANT: Duclert,

APPLICANT: Glordano, J.Y.

TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.

PATENT FILE REFERENCE: 59-US2.REG

CURRENT FILING DATE: 2000-02-24

PRIOR PILING DATE: 1999-02-26

NUMBER OF SEQ ID NOS: 36681

SOFTWARE: Patent.pm

SEQ ID NO 32786

LENGTH: 95

LENGTH: 95

LENGTH: 95

LENGTH: 95

LENGTH: 95
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                                                                                                                                                                                                                                                                                                                                                                                                                                    1 TAGTAGAGACGAGGTTTCACCGAATTAGCCAGGATGGTCTCGATCTCCTGACCTTGTGAT 60
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Patent No. 6783961
GENERAL INFORMATION:
APPLICANT: Dunas Milne Edwards, J.B.
APPLICANT: Dunas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Duclert, A.
APPLICANT: Giordano, J.Y.
FAPPLICANT: Giordano, J.Y.
FAPPLICANT: Giordano, J.Y.
FAPPLICANT: Giordano, J.Y.
CURRENT PREFIERCY: 59.US2.REG
CURRENT PELIGNON NUMBER: US/09/513,999C
CURRENT PELING DATE: 2000-02-24
FRIOR APPLICATION NUMBER: US 60/122,487
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                                                                                                                                                                                                                                                                                                                                            8; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    61 CTGCCCGCCTTGGCCTCCCAAAGTGCTGGGATTA 94
PRIOR APPLICATION NUMBER: US 60/122,487 PRIOR FILING DATE: 1999-02-26 NUMBER OF SEQ ID NOS: 36681 SOFTWARE: Patent.pm
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ORGANISM: Homo sapiens
US-09-513-999C-32786
                                                                                                                                                                       ; TYPE: DNA
; ORGANISM: Homo'sapiens
US-09-513-999C-28026
                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Dunas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Duclert, A.
APPLICANT: Glocdano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
Patent No. 6783961
FILE REPERENCE: 59.052.786
FILE REPERENCE: 59.052.786
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SEQ ID NO 27645
LENGTH: 90
                                                                                                                                                                                                                                                                                                                                                    96 TITITIGIATITITAGIAGAGACAGGGITICAGCAIGITGGCCAGGCIGGICCAAACIC 37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     90 TITITITITITITIAGIAGAGAGGGGTTTCACCGTGTTAGCCAGGATGGTCCCATCTC 31
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Sequence 19069, Application US/09513999C

Sequence 19069, Application US/09513999C

Sequence 19069, Application US/09513999C

GRNERAL INFORMATION:

APPLICANT: Dumas Milne Edwards, J.B.

APPLICANT: Duclert, A.

APPLICANT: Glordano, J.Y.

TILE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.

Patent No. 6783961

FILE REFERENCE: 59.US2.REG
                                                                                                                                                                                                                                                                             Gaps
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Pred. No. 0.045;
1; Mismatches 7; Indels
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3.4%; Score 80; DB 1; Length 100;
Best Local Similarity 89.6%; Pred. No. 0.035;
Matches 86; Conservative 0; Mismatches 10; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                   2310 CIGACCICGIGATCCGCCCACCTCGGCCTCCCAAAG 2345
                                                                                                                                                                                                                                                                                                                                                                                                                                                              36 CTGACCTTGTGATCCATCCACCTCGGCCTCCCAAAG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-513-999C-27645/c
; Sequence 27645, Application US/09513999C
; Patent No. 678361
; GENERAL INFORMATION:
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Best Local Similarity 91.1%;
Matches 82; Conservative
PRIOR FILING DATE: 1999-02-26
                    NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 15118
LENGTH: 100
TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: m=a or c
US-09-513-999C-27645
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ORGANISM: Homo sapiens
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Sequence 49, Application US/09073567

APPLICANT: Standong Chen

APPLICANT: Sudhir Agrawal

APPLICANT: Ruiwen Zhang

TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES

NUMBER OF SEQUENCES: 49

CORRESPONDENCE ADDRESS:

ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff

STREET: 300 South Wacker Drive, 32nd Floor

CITY: Chicago
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                                                                                                                                                                                                                                                                                    Length 86;
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                                                                                                                                                                                                                                                                                                                               5; Indels
                                                                                                                                                                                                                                                                               Score 78; DB 1;
Pred. No. 0.049;
0; Mismatches
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 19069
LENGTH: 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2320 GATCCGCCCACCTCGGCCTCCCAAAG 2345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                26 GATCCTCCCACCTCGGCCTCACAAAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 37,147
TELEPHONE: (312) 913-0001
TELEPAX: (312) 913-0001
INFORMATION FOR SEQ ID NO: 49:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: nucleic acid HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                 Query Match 3.3%;
Best Local Similarity 94.2%;
Matches 81; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       73 base pairs
                                                                                                                                                                                                              ; ORGANISM: Homo sapiens
US-09-513-999C-19069
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: both
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: UN
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Batent No. 6783961

GENERAL INPORMATION:
APPLICANT: Dunas Milne Edwards, J.B.
APPLICANT: Dunas Milne Edwards, J.B.
APPLICANT: Dunclert, A.
APPLICANT: Duclert, A.
TILE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE REPRENCE: 59.U32.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR PILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOUTHWARE: PARENTE: NOS: 36681
SOUTHWARE: PARENTE: DATE: D
                                                                                                                                                                                                                                                                                                                                                                                                        US-00-513-999C-17764/C

Sequence 17764, Application US/09513999C

Patent No. 6783961

APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
PILE REFERENCE: 59.US2.REG

CURRENT FILING DATE: 2000-02-24

PRIOR APPLICATION NUMBER: US 60/122,487

PRIOR PLING DATE: 1999-02-26

NUMBER OF SEQ.ID NOS: 36681

SEQ ID NO 17764

LENGTH: 76
665 AGGTACATCTGTGAGTGAGAACAGGTGTCACCTTGAAGGTGGGAGTGATCAAAAGGACCT 724
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           76 TGTTAGCCAGGATGGTCTCGATCTCCTGACCTCGTGATCCGCCCTCCTCAGCCTCCCAAA 17
                                             13 AGGTACATCTGTGAGTGAGAGAGAGTGTCACCTTGAAGGTGGGGGGTGATCAAAAGGACCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 76;
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3.1%; Score 72.8; DB 1;
Best Local Similarity 97.4%; Pred. No. 0.11;
Matches 74; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2345 GTGCTGGGATTACAGG 2360
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; ORGANISM: Homo sapiens
US-09-513-999C-17764
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US-09-513-999C-16592
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LENGTH: 84
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2248 AATTTTTGTACTTTTAGTAGAGACAGGGTTTCACCGTGTTAGCCAGGATGGTCTCGATC 2307

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CTCGATCTCCT 2311
                                                                                                                  RESULT 12
US-09-513-999C-28108
i Sequence 28108, Application US/09513999C
j Patent No. 6783961
j GENERAL INFORMATION:
i APPLICANT: Dunas Milne Edwards, J.B.
i APPLICANT: Glordano, J.Y.
i TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
j Patent No. 6783961
j FILE REPERENCE: 59.9US2.REG
cURRENT FILING DATE: 2000-02-24
j PRIOR PELICATION NUMBER: US 60/122,487
j RIOR FILING DATE: 1999-02-26
j NUMBER OF SEQ ID NOS: 36681
j SOFTWARE: Patent.pm
j SEQ ID NO 28108
j LENGTH: 79
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0; Gaps
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Best Local Similarity 85.7%; Pred. No. 0.15;
Matches 78; Conservative 0; Mismatches 13;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 71; DB 1
Pred. No. 0.14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
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% Sequence 15415, Application US/09513999C
% Patent No. 6783961
% GRNEAAL INFORMATION:
% APPLICANT: Dumas Milne Edwards, J.B.
                                              2308 TCCTGACCTCGTGATCCGCCC 2328
                                                               62 TCCTGACCTCGTGATCCACCC 82
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Best Local Similarity 93.7%;
Matches 74; Conservative
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CRGANISM: Homo sapiens
US-09-513-999C-15415
                                                                                                                                                                                                                                                                                                                                                                                                                  ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-513-999C-28108
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Page 10

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US-08-340-426D-91
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US-08-454-557C-91
Sequence 91, Application US/08454557C
Fatent No. 5830670
GENERAL INFORMATION:
APPLICANT de la Monte, Suzanne
APPLICANT Hands, Jack R.
TITLE OF INVENTION: Neural Thread Protein Gene Expression and Detection
TITLE OF INVENTION: Net all Thread Protein Gene Expression and Detection
TITLE OF INVENTION: Of Alzheimer's Disease
NUMBER OF SEQUENCES: 121
CORRESPONDENCE ADDRESS: 121
CORRESPONDENCE ADDRESS: 121
CORRESPONDENCE ADDRESS: 121
CORRESPONDENCE ADDRESS: 121
CONDETT: 1100 New York Avenue, Suite 600
STATE: 100.
COMPUTER: IBM FOC Compatible
COMPUTER: IBM FOC Compatible
COMPUTER: IBM FOC Compatible
COMPUTER: 18M FOC COMPATIBLE
COMPUTER: 18M FOC COMPATIBLE
COMPUTER: 18M FOC COMPATIBLE
CORPUTER: 18M FOC COMPATIBLE
COMPATIBLE
COMPATIBLE
CORPUTER
COMPATIBLE
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1 TTTTTGTATTTTTAGTAGACGGGTTTCACCATGTTGGTCAGGCTGGTCTTGAACTCC 60
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 ATTITICATAGAGACATGGTTTCACCATGTTAGCCATGATGGTCTCGATCTCCTGACCTC 60
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                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INCOMMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Jobert, S.
APPLICANT: Glordanch, J.Y.
ITILE OF INVENTION: ESTS and Encoded Human Proteins.
FILE REFERENCE: GENSET.054FR2
CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 19335
SOFTWARE: Parent.pm
SEQ ID NO 12767
LENGTH: 85
                                                                           2311 TGACCICGIGATCCGCCCACCICGGCCTCCC 2341
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REGISTRATION NUMBER: 36,203
REFERENCE/DOCKET NUMBER: 0609.3840003
FELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2318 GIGATCCGCCCACCTCGGCCTCCCA 2342
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US-09-621-976-12767
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: de la Monte, Suzanne
APPLICANT: de la Monte, Suzanne
APPLICANT: Wands, Jack
APPLICANT: Wands, Jack
TITLE OF INVENTION: Neural Thread Protein Gene Expression and Detection
TITLE OF INVENTION: of Alzheimer's Disease
NUMBER OF SEQUENCES: 121
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox P.L.L.C.
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                                                                                                                                                                                               Score 65.4; DB 1; Length 84;
Pred. No. 0.32;
0; Mismatches 11; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: Sterne, Kessler, Goldstein & Fox P.L.L.C. STREET: 1100 New York Avenue, Suite 600 STATE: D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FURM:
MEDIUM TYEE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
SUFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/340,426D
FILING DATE: 14-NOV-1994
CLASSIFICATION: 435
ATTORNEY AGENT INFORMATION:
NAME: Ludwig, Steven R:
REFERENCE/POCKET NUMBER: 0609.3840002
FELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OF SECONO TELECOMMUNICATION OF SECONO TELECOMMUNICATION:
TELECOMMUNICATION: 91:
SEQUENCE CHARACTERISTICS:
TWATH: 84 base pairs
                                                                                                                                                                                                                                                                                                   2282 CCGTGTTAGCCAGGATGGTCTCGATCTCCTGACCTCGTGA
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; Sequence 91, Application US/08340426D
; Patent No. 5948634
; INFORMATION FOR SEQ ID NO:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 84 base pairs
; TYPE: nucleic acid
; STRANDENBESS: both
; TOPOLOGY: both
US-08-454-557C-91
                                                                                                                                                                           Query Match
Best Local Similarity 86...
Best Local Similarity 86...
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Matches 72; Conservative
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COUNTRY: U.S.A.
ZIP: 20005-3934
COMPUTER READABLE FORM:
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STRANDEDNESS: both
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             Sequence 91, Application US/08450673C

Sequence 91, Application US/08450673C

Patent No. 594888

GENERAL INFORMATION:
APPLICANT: de la Monte, Suzanne
APPLICANT: de la Monte, Suzanne
APPLICANT: Wands, Jack R.
ITTLE OF INVENTION: of Alzheimer's Disease
NUMBER OF SEQUENCES: 121
CORRESPONDENCES: 122
CORRESPONDENCES: 122
CORRESPONDENCES: 121
CORRESPONDENCES: 122
CONTEXT: DC.
STREET: 1100 New York Avenue, Suite 600
CITY: Washington
STATE: DC.
COUNTRY: U.S.A.
ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: DC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/450,673C
FILING DATE: 30-MAY-1995
ATTORNEY/AGENT INFORMATION:
NAME: Linding Carenar DR.
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GENERAL INFORMATION:
APPLICANT: de la Monte, Suzanne
APPLICANT: de la Monte, Suzanne
APPLICANT: Wands, Jack R.
TITLE OF INVENTION: Neural Thread Protein Gene Expression and
TITLE OF INVENTION: Detection of Alzheimer's Disease
NUMBER OF SEQUENCES: 121
CORRESPONDENCE ADDRESS:
ADDRESSEE: Steine, Kessler, Goldstein & Fox P.L.L.C.
STREET: 1100 New York Avenue, Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2.8%; Score 65.4; DB 1; Length 84; 86.7%; Pred. No. 0.32;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Ludwig, Steven R.
REGISTRATION NUMBER: 36,203
REFERENCE/DOCKET NUMBER: 0609.
TELECOMMUNICATION INPORMATION:
TELEPAX: (202) 371-2500
INPORMATION FOR SEQ ID NO: 91:
SEQUENCE CHARACTERISTICS:
LENGTH: 84 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 2.8%
Best Local Similarity 86.7%
Matches 72; Conservative
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ZIP: 20005-3934
COMPUTER READABLE FORM:
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STATE: D.C.
COUNTRY: U.S.A.
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US-08-450-673C-91
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2282 CCGTGTTAGCCAGGATGGTCTCGATCTCCTGACCTCGTGATCCGCCCACCTCGGCCTCCC 2341
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US-09-513-999C-16016

Sequence 16016, Application US/09513999C

Patent No. 6783961

GENERAL INFORMATION:

APPLICANT: Duclart, A.

APPLICANT: Duclart, A.

TITLE OF INVENTION:

Patent No. 6783961

CURRENT APPLICATION NUMBER: US/09/513,999C

CURRENT FILING DATE: 2000-02-24

PRIOR FILING DATE: 1999-02-26

NUMBER: OF SEQ ID NOS: 36681

SEQ ID NO 16016

LENGTH: B.

LENGTH: B.
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Pred. No. 0.32;
0; Mismatches 11; Indels
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US95/17111A
FILING DATE: CLASSIFICATION:
PRIOR APPLICATION NUMBER: 08/340,426
FILING DATE: 14-NOV-1994
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
NAME: Lidwing Steven R.
REGISTRATION NUMBER: 36,203
REFERENCE/DOCKET NUMBER: 0609.3840002
TELEPRAK: (202) 371-2540
INFORMATION EN REQ ID NO: 91:
SEQUENCE CHARACTERISTICS:
LENGTH: 84 base pairs
TYPE: nucleic acid
STRANDBNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2342 AAAGTGCTGGGATTACAGGCATG 2364
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Best Local Similarity 86.7%;
Matches 72; Conservative (
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// LOCATION: 48
// OTHER INFORMATION: r=a or
US-09-513-999C-16016
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OTHER INFORMATION: n=a,
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LOCATION: 8
OTHER INFORMATION: n=a,
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ORGANISM: Homo sapiens
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PCT-US95-17111A-91
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2253 TITGIACTITIAGIAGACAGGGTITCACCGTGTTAGCCAGGATGGTCTCGATCTCCTG 2312
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                                                                                                                                                                                                                                                                                                     Sequence 17000, Application US/09513999C
; Sequence 17000, Application US/09513999C
; Patent No. 6783961
; GENERAL INFORMATION:
; APPLICANT: Dumas Milne Edwards, J.B.
; APPLICANT: Duclert, A.
; APPLICANT: Glordano, J.Y.
; TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
; Patent No. 678399C
; FILE REPRENCE: 59, US2. REG
; CURRENT FILING DATE: 2000-02-24
; PRIOR PPLICATION NUMBER: US 60/122,487
; PRIOR FILING DATE: 1999-02-26
; NUMBER OF SEQ ID NOS: 36681
; SEQ ID NO 17000
; SEQ ID NO 17000
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                                                                                                                            2 TATTINNITIAAGIAGAGACAGGGTTTCATCGTGTTTGCCAGGATGRTCTCAAACTCCTG 61
                                                Gaps
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2.6%; Score 62.2; DB 1; Length 81;
84.8%; Pred. No. 0.52;
tive 1; Mismatches 11; Indels
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Pred. No. 2.5;
0; Mismatches 10; Indels
                                           11; Indels
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Patent No. 6639063
GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Jobert, S.
APPLICANT: Glordano, J.Y.
TITLE OF INVENTION: ESTS and Encoded Human Proteins.
FILE REFERENCE: GENSET.054PR2
CURRENT APPLICATION NUMBER: US/09/621,976
CURRENT PILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 19335
SOCTWARRE: Patent.pm
SEQ ID NO 7999
LENGTH: 68
                                                                                                                                                                      2313 ACCTCGTGATCCGCCCACC 2331
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                                         Matches 67; Conservative
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Matches 58; Conservative
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Best Local Similarity 85.3
Matches 58; Conservative
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US-09-621-976-7999
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ORGANISM: Homo sapiens
                     Best Local Similarity
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    Query Match
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APPLICANT: Van Ommen, Garrit J.B.
APPLICANT: Petrij-Bosch, Anne
APPLICANT: Petrij-Bosch, Anne
APPLICANT: Bakker, Egbert
APPLICANT: Bakker, Egbert
APPLICANT: Bakker, Egbert
APPLICANT: Devilee, Peter
TITLE OF INVENTION: A diagnostic test kit for determining a predisposition
TITLE OF INVENTION: for breast and ovarian cancer, materials and methods
TITLE OF INVENTION: for such determination
FILE REPERENCE: 294-78
FILE REPERENCE: 294-78
CURRENT PILING DATE: 2001-06-11
PRIOR FILING DATE: 1998-06-03
PRIOR APPLICATION NUMBER: EP 97201700.8
PRIOR PLING DATE: 1997-06-04
NUMBER OF SEQ ID NOS: 23
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 23
LINGUIL OF ALCOME ADDITIONAL ADDITIONAL
2189 TCTCCTGCCTCAGCCTCCCAATTAGCTTGGCCTACAGTCATCTGCCACCACACCTGGCTA 2248
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Sequence 15107, Application US/09513999C

Patent No. 6783961

GENERAL INFORMATION:
APPLICANT: Dunas Milne Edwards, J.B.
APPLICANT: Dunas Milne Edwards, J.B.
APPLICANT: Dunalert, A.
APPLICANT: Discrepance 100 of 100 o
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                                                                                                                          68 TCTCCTGCCTCAGCCTCCCGAGTAGCTGGGACTACAGGCACCCGCCACCGCGCCCAC
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Pred. No. 2.8;
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US-09-445-174B-23
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Best Local Similarity 91.5%; Pred. No. 2.9;
Matches 54; Conservative 0; Mismatches 5
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; Sequence 23, Application US/09445174B
; Patent No. 6733966
; GENERAL INFORMATION:
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85.1%;
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Best Local Similarity
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                                                                                                                                            2251 TTTTTGTACTTTTAGTAGAGAGAGGTTTCACCGTGTTAGCCAGGATGGTCTCGATCTCC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:

APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Duclert, A.
APPLICANT: Duclert, A.
APPLICANT: Diordano, J.Y.
TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
PILE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
FILE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
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Sequence 1171, Application US/09443199C

Parent No. 667044

GRNERAL INFORMATION:

APPLICANT: Shimkets, Richard A.

APPLICANT: Leach, Martin

TITLE OF INVENTION: Nucleic Acids Containing Single Nucleotide

TITLE OF INVENTION: Nucleic Acids Containing Single Nucleotide

TITLE OF INVENTION: Polymorphisms and Methods of Use Thereof

FILE REFERENCE: 15966-534A

CURRENT FILING DATE: 1999-11-16

PRIOR APPLICATION NUMBER: 60/109,024

PRIOR PLICATION TOWNER: 1999-11-17

PRIOR PLICATION TOWNER: 1999-11-17
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OTHER INFORMATION: 1 of 2 allelic variants (1172 is other entry)
NAME/KEY: misc_feature
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        10; Indels
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SOFTWARE: CuraGen Patent Formatter Version 0.9
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Pred. No. 3.2;
0; Mismatches
    Mismatches
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Best Local Similarity 87.3%;
Matches 55; Conservative (
57; Conservative
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ORGANISM: Homo sapiens
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US-09-513-999C-19798
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LENGTH: 64
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SEQ ID NO 1171
LENGTH: 51
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GENERAL INFORMATION:
APPLICANT: Cohen, Daniel
APPLICANT: Cohen, Daniel
APPLICANT: Bluemerlead, marta
APPLICANT: Chumakov, Ilya
TITLE OF INVENTION: Biallelic markers for use in constructing a high density...
FILE REFERENCE: GENSET. O20CP1
CURRENT APPLICATION NUMBER: US/09/422,978
CURRENT FILING DATE: 1999-10-20
EARLIER PEPLICATION NUMBER: US 09/298,850
EARLIER FILING DATE: 1999-04-21
EARLIER PLING DATE: 1999-11-23
EARLIER PLING DATE: 1998-04-21
EARLIER PLING DATE: 1998-04-21
EARLIER PLING DATE: 1998-04-21
EARLIER PLING DATE: 1998-04-21
NUMBER OF SEQ ID NOS: 11796
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OTHER INFORMATION: 2 of 2 allelic variants (1171 is other entry)
NAME/EN: misc_feature
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GENERAL INFORMATION:
APPLICANT: Shimkets, Richard A.
APPLICANT: Leach, Martin
TITLE OF INVENTION: Nucleic Acids Containing Single Nucleotide
TITLE OF INVENTION: Polymorphisms and Methods of Use Thereof
TITLE OF INVENTION: Polymorphisms and Methods of Use Thereof
TITLE OF INVENTION: 15966-534A
CURRENT APPLICATION NUMBER: 105/09/443,199C
CURRENT FILING DATE: 1999-11-16
PRIOR FILING DATE: 1998-11-17
NUMBER OF SEQ ID NOS: 1272
SOFTWARE: Curaden Patent Formatter Version 0.9
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                                                                       Length 51;
                                                                                                                     1; Indels
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                                                                    Score 48.4; DB 1;
Pred. No. 4.4;
0; Mismatches 1;
; OTHER INFORMATION: Accession number cg43129081
US-09-443-199C-1171
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; OTHER INFORMATION: Accession number cg43129081
US-09-443-199C-1172
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; Patent No. 6537751
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96.0%;
                                                                  Query Match 2.0%;
Best Local Similarity 98.0%;
Matches 49; Conservative
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Best Local Similarity 96.09
Matches 48; Conservative
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ORGANISM: Homo sapiens
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ORGANISM: Homo Sapiens
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US-09-443-199C-1172
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LENGTH: 51
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US-09-513-999C-33702, Application US/09513999C

Sequence 33702, Application US/09513999C

Sequence 33702, Application US/09513999C

GENERAL INFORMATION:

APPLICANT: Dumas Milne Edwards, J.B.

APPLICANT: Duclert, A.

APPLICANT: Glordano, J.Y.

TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.

FILE REFERENCE: 59.US2.REG

CURRENT APPLICATION NUMBER: US/09/513,999C

CURRENT FILING DATE: 2000-02-24

PRIOR FILING DATE: 1999-02-26
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Sequence 671, Application US/09443199C

Retent No. 6670464

GENERAL INFORMATION:

APPLICANT: Shinkers Ratin

TITLE OF INVENTION: Nucleic Acids Containing Single Nucleotide

TITLE OF INVENTION: Nolleic Acids Containing Single Nucleotide

TITLE OF INVENTION: Polymorphisms and Methods of Use Thereof

TITLE OF INVENTION NUMBER: US/09/443,199C

CURRENT APPLICATION NUMBER: US/09/443,199C

CURRENT FILING DATE: 1999-11-16

PRIOR FILING DATE: 1999-11-17

NUMBER OF SEQ ID NOS: 1272

SOFTWARE: CURRENT CURRENT PRICE CURRENT FORMATHER VERSION 0.9

SEQ ID NO 671

LENGTH: 51
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LOCATION: (26)...(0)
OTHER INFORMATION: 1 of 2 allelic variants (672 is other entry)
NAME/KEY: misc_feature
LOCATION: (0)...(0)
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                                                                                                                             2.0%; Score 46.6; DB 1; Length 47; 97.9%; Pred. No. 5.9; ive 1; Mismatches 0; Indels
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NAME/KEY: allele
LOCATION: 24
J. DICATION: 24
J. OTHER INFORMATION: 99-4582-359 : polymorphic base G or US-09-422-978-3882
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87.7%; Pred. No. 6.4;
iive 0; Mismatches 7;
                                                                                                                        Query Match
Best Local Similarity 97.93
Matches 46; Conservative
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Best Local Similarity 87.7
Matches 50; Conservative
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SOFTWARE: Patent.pm
SEQ ID NO 33702
LENGTH: 57
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ORGANISM: Homo sapiens
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US-09-513-999C-33702
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US-09-443-199C-671/c
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Sequence 659, Application US/09641638

Patent No. 6432649

GENERAL INFORMATION

APPLICANT: Blumenfeld, Marta

APPLICANT: Chumakov, Ilya

TITLE OF INVENTION: GENETI SIQUE ENSET.051CP1

TITLE OF INVENTION: GENETI 051CP1

CURRENT APPLICANTON NUMBER: US/09/641, 638

CURRENT APPLICANTON NUMBER: US 09/502,330

PRIOR APPLICATION NUMBER: US 09/502,330

PRIOR APPLICATION NUMBER: US 09/502,330

PRIOR APPLICATION NUMBER: US 09/275,267

PRIOR APPLICATION NUMBER: US 09/275,267

PRIOR FILING DATE: 1999-05-07

PRIOR FILING DATE: 1999-03-23

PRIOR FILING DATE: 1999-02-12

NUMBER OF SEQ ID NOS: 1304
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                                                                                                                                                                                                                                                                                                                                                        | Sequence 1125, Application US/09443199C |
| Patent No. 6670464 |
| GENERAL INFORMATION |
| APPLICANT: Shimkets, Richard A. |
| APPLICANT: Leach, Martin |
| TITLE OF INVENTION: Nucleic Acids Containing Single Nucleotide |
| TITLE OF INVENTION: Polymorphisms and Methods of Use Thereof |
| TITLE OF INVENTION: Polymorphisms and Methods of Use Thereof |
| FILE REPERENCE: 15966-34A |
| CURRENT APPLICATION NUMBER: US/09/443,199C |
| PRIOR PILING DATE: 1998-11-16 |
| PRIOR FILING DATE: 1998-11-17 |
| NUMBER OF SEQ ID NOS: 1272 |
| SOFTWARE: Curaden Patent Formatter Version 0.9 |
| SEQ ID NO 1125
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Homo sapiens
FEATURE:
NAME/KEY: misc feature
LOCATION: (26)...(0)
OTHER INFORMATION: 1 of 2 allelic variants (1126 is other entry)
NAME/KEY: misc feature
LOCATION: (0)...(0)
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                                                                                   Length 51;
                                                                                   DB 1;
; OTHER INFORMATION: Accession number cg42924993
US-09-443-199C-671
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OTHER INFORMATION: Accession number cg42894694
                                                                             Score 43; DB 1; Pred. No. 9.8; 0; Mismatches
                                                                             1.8%;
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                                                                             Query Match
Best Local Similarity
Matches 46; Conserv
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Page 15

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APPLICANT: Bougueleret, Lydie
APPLICANT: Chumakov, Ilya
TITLE OF INVENITON: BIALLELIC MARKERS DERIVED FROM GENOMIC REGIONS CARRYING
TITLE OF INVENITON: GENES INVOLVED IN ARACHIDONIC ACID METABOLISM
FILE REPREBACE: GEN-T114XC2DI
CURRENT APPLICATION UNMBER: US/10/170,097
CURRENT APPLICATION NUMBER: US 09/641,638
PRIOR PILING DATE: 2000-08-16
PRIOR FILING DATE: 2000-02-11
PRIOR FILING DATE: 1999-03-23
PRIOR PILING DATE: 1999-03-23
PRIOR FILING DATE: 1999-02-12
NUMBER OF SEQ ID NOS: 1304
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                                                                                                                                                                                                                                                                                                                                     2274 GGGTTTCACCGTGTTAGCCAGGATGGTCTCGATCTCCTGACCT 2316
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COTHER INFORMATION: 10-520-256 : polymorphic base C or T
US-10-170-097-659
                                                                                                                                  NAME/KEY: allele
COATION: 24

US-09-641-638-659
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                                                                                                                                                                                                                                               Score 42.6; DE
Pred. No. 11;
1; Mismatches
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US-09-422-978-646/c
US-09-422-978-646/c
; Sequence 646, Application US/09422978
; Patent No. 6537751
; GENERAL INFORMATION:
; APPLICANT: Cohen, Daniel
; APPLICANT: Chumanfeld, Marta
; APPLICANT: Chumakov, Ilya
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 659, Application US/10170097
Patent No. 6794143
                                                                                                                                                                                                                                                 1.8%;
                                                                                                                                                                                                                                          Query Match
Best Local Similarity 97.7%
Matches 42; Conservative
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Matches 42; Conservative
                                                                   TYPE: DNA
ORGANISM: Homo Sapiens
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ORGANISM: Homo Sapiens
    Patent.pm
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SOFTWARE: Pa
SEQ ID NO 659
LENGTH: 47
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LENGTH: 47
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TITLE OF INVENTION: Biallelic markers for use in constructing a high density...
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| Patent No. 6783961
| GENERAL INPORMATION:
| APPLICANT: Dumas Milne Edwards, J.B.
| APPLICANT: Dumas Milne Edwards, J.B.
| APPLICANT: Duclert, A.
| TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
| PATENT REPERENCE: 59. US2.REG
| PATENT PAPLICATION NUMBER: US/09/513,999C
| CURRENT APPLICATION NUMBER: US 60/122,487
| PRIOR PAPLICATION NUMBER: US 60/122,487
| PRIOR FILING DATE: 1999-02-26
| NUMBER OF SEQ ID NOS: 36681
| SOFTWARE: Patent.pm
| SEQ ID NO 26492
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; OTHER INFORMATION: 99-1602-200 : polymorphic base G or C
US-09-422-978-646
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Best Local Similarity 93.5%; Pred. No. 11;
Matches 43; Conservative 1; Mismatches
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86.5%; Pred. No. 12;
cive 1; Mismatches
                                          FILE REFERENCE: GENSET.020CP1
CURRENT APPLICATION NUMBER: US/09/422,978
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US 09/298,850
EARLIER FILING DATE: 1999-04-21
EARLIER FILING DATE: 1999-04-21
EARLIER PLILNG DATE: 1998-11-23
EARLIER PLILNG DATE: 1998-11-23
EARLIER FILING DATE: 1998-11-35
EARLIER FILING DATE: 1998-04-21
EARLIER FILING DATE: 1998-04-21
SEQ. ID NO 646
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10.09-443-199C-672/C

10.09-431-199C-672/C

10.09-491-109C

10.09-4
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US-09-513-999C-26492
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Best Local Similarity 86.5
Matches 45; Conservative
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ORGANISM: Homo Sapiens
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JS-09-443-199C-911
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Patent No. 6670464

GENERAL INFORMATION:
APPLICANT: Shimkets, Richard A.
APPLICANT: Leach, Martin
TITLE OF INVENTION: Nucleic Acids Containing Single Nucleotide
TITLE OF INVENTION: 1904-9140, 1995
CURRENT APPLICATION NUMBER: 1090-11-16
PRIOR PLILING DATE: 1999-11-16
PRIOR FILING DATE: 1999-11-17
NUMBER OF SEQ ID NOS: 1272
SOFTWARE CURRENT PATENT PATENT
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LOCATION: (26)...(0)

OTHER INFORMATION: 2 of 2 allelic variants (671 is other entry)

NAME/KEY: misc feature

LOCATION: (0)...(0)

OTHER INFORMATION: Accession number cg42924993

US-09-443-199C-672
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LOCATION: (26)...(0)
OTHER INFORMATION: 1 of 2 allelic variants (914 is other entry)
NAME/KEY: misc_feature
LOCATION: (0)...(0)
OTHER INFORMATION: Accession number cg43972482
                INVENTION: Nucleic Acids Containing Single Nucleotide INVENTION: Polymorphisms and Methods of Use Thereof
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1.7%; Score 41.4; DB 1; Length 51;
88.2%; Pred. No. 12;
tive 0; Mismatches 6; Indels
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TITLE OF INVENTION: Nucleic Acids Containing Singl.
TITLE OF INVENTION: Polymorphisms and Methods of
FILE REFERENCE: 15966-534A
CURRENT APPLICATION NUMBER: US/09/443,199C
CURRENT FILING DATE: 1999-11-16
PRIOR APPLICATION NUMBER: 60/109,024
PRIOR PILING DATE: 1998-11-17
NUMBER OF FILING DATE: 1998-11-17
SOFTWARE: CuraGen Patent Formatter Version 0.9
SEQ ID NO 672
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US-09-443-199C-1126
Sequence 1126, Application US/09443199C
; Pacent No. 6670464
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88.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 88.2<sup>3</sup>
Matches 45; Conservative
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
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Best Local Similarity
Matches 45; Conserva
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US-09-443-199C-913
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LENGTH: 51
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APPLICANT: Shimkete, Richard A.
APPLICANT: Leach, Martin
TITLE OF INVENTION: Nucleic Acids Containing Single Nucleotide
TITLE OF INVENTION: POLYMorphisms and Methods of Use Thereof
TITLE OF INVENTION: POLYMorphisms and Methods of Use Thereof
TITLE OF INVENTION: POLYMorphisms and Methods of Use Thereof
CURRENT APPLICATION NUMBER: US/09/443,199C
CURRENT FILING DATE: 1999-11-16
PRIOR PILING DATE: 1998-11-17
NUMBER OF SEQ ID NOS: 1272
SOFTWARE: CuraGen Patent Formatter Version 0.9
SEQ ID NO 911
LENGTH: 51
                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA

OKGANISM: Homo sapiens
FEATURE:
NAMB/KEX: misc feature
LOCATION: (26)...(0)
OTHER INFORMATION: 2 of 2 allelic variants (1125 is other entry)
LOCATION: (0)...(0)
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OTHER INFORMATION: 1 of 2 allelic variants (912 is other entry)
NAME/KEY: misc_feature
                    APPLICANT: LEACH, MARKING TITLE OF INVENTION: MACHINE ACIDS CONTAINING SINGLE Nucleotide TITLE OF INVENTION: Nucleic Acids Containing Single Nucleotide FITLE OF INVENTION: Polymorphisms and Methods of Use Thereof FILE REFERENCE: 1596-534A
CURRENT APPLICATION NUMBER: US/09/443,199C
CURRENT FILING DATE: 1999-11-16
PRIOR APPLICATION NUMBER: 60/109,024
PRIOR FILING DATE: 1998-11-17
NUMBER OF SEQ ID NOS: 1272
SEQ ID NO 126
LENGTH: 51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 1.7%; Score 40.4; DB 1; Length 51; Best Local Similarity 97.6%; Pred. No. 14; Matches 41; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 51;
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    LOCATION: (0)...(0)
    OTHER INFORMATION: Accession number cg42894694
US-09-443-199C-1126

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OTHER INFORMATION: Accession number cg43971764
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.7%; Score 41.4; Dilarity 88.2%; Pred. No. 12; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 911, Application US/09443199C Patent No. 6670464 GENERAL INFORMATION:
Richard A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: misc_feature
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 45; Conserv
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GENERAL INFORMATION:

Wed Jan 26 12:47:30 2005

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US-09-513-999C-18997/C

Sequence 18997, Application US/09513999C

Patent No. 678361

APPLICANT: Duclert, A.

APPLICANT: Giordano, J.Y.

TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.

PATENTE SPERENCE: 59.US2.REG

FILE REFERENCE: 59.US2.REG

CURRENT APPLICATION NUMBER: US/09/513,999C

CURRENT FILING DATE: 2000-02-24

PRIOR FILING DATE: 10999-02-26

NUMBER OF SEQ 1D NOS: 36681

SOFTWARE: PATENT DATE: DATE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Gaps
              Sequence 294, Application US/09443199C

Patent No. 6670464

GENERAL INFORMATION:

APPLICANT: Shimkers, Richard A.

APPLICANT: Shimkers, Richard A.

TITLE OF INVENTION: Nucleic Acids Containing Single Nucleotide

TITLE OF INVENTION: Polymorphisms and Methods of Use Thereof

TILE REPREBUCE: 15966-534A

CURRENT APPLICATION NUMBER: US/09/443,199C

CURRENT APPLICATION NUMBER: 0999-11-16

PRIOR APPLICATION NUMBER: 60/109,024

PRIOR FILING DATE: 1998-11-17

NUMBER OF SEQ ID NOS: 1272

SOFTWARE: Curaden Patent Formatter Version 0.9

SEQ ID NO 914
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION: (26)...(0)
OTHER INFORMATION: 2 of 2 allelic variants (913 is other entry)
NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
1.7%; Score 39.8; DB 1; Length 51;
Best Local Similarity 86.3%; Pred. No. 16;
Matches 44; Conservative 0; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1.7%; Score 39.8; DB 1; Length 51;
36.3%; Pred. No. 16;
Lve 0; Mismatches 7; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LOCATION: (0)...(0)
OTHER INFORMATION: Accession number cg43972482
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 1.7%;
Best Local Similarity 86.3%;
Matches 44; Conservative
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ORGANISM: Homo sapiens
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US-09-513-999C-18997
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US-09-443-199C-912
US-09-443-199C-914
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LENGTH: 51
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OTHER INFORMATION: Description of Artificial Sequence:Oligonucleotide
OTHER INFORMATION: that is complementary to a portion of the human
OTHER INFORMATION: MDM2 mRNA
APPLICANT: Shimkets, Richard A.
APPLICANT: Leach, Martin
TITLE OF INVENTION: Nucleic Acids Containing Single Nucleotide
TITLE OF INVENTION: Nucleic Acids Containing Single Nucleotide
TITLE OF INVENTION: POLYMOTPHISMS and Methods of Use Thereof
FILE REFERENCE: 15966-534A
CURRENT APPLICATION NUMBER: US/09/443,199C
CURRENT FILING DATE: 1999-11-16
PRIOR APPLICATION NUMBER: 60/109,024
PRIOR PILING DATE: 1998-11-17
NUMBER OF SEQ ID NOS: 1272
SOFTWARE: Curaden Patent Formatter Version 0.9
SEQ ID NO 912
LENGTH: 51
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1.6%; Score 38.8; DB 1; Length 51;
Best Local Similarity 95.2%; Pred. No. 18;
Matches 40; Conservative 0; Mismatches 2; Indels
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Patent No. 6383752;
GENERAL INFORMATION:
FAPPLICANT: APPLICANT: Randimalla, Ekambar R.
TITLE OF INVENTION: Peetdo-Cyclic Oligonucleobases;
FILE REFERENCE: 99,128-B;
CURRENT FILING DATE: 2000-03-31;
PRIOR FILING DATE: 1999-03-31;
PRIOR FILING DATE: 1999-03-31;
PRIOR FILING DATE: 2000-01-05;
NUMBER OF SEQ ID NOS: 26
SOSTWARE: PatentIn Ver. 2.0
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LOCATION: (0)...(0)

CTHER INFORMATION: Accession number cg43971764

US-09-443-199C-912
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97.5%; Pred. No. 20;
tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 97.5'
                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: misc_feature
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                                                                                                                                                                                                                                                                                                              1 TCCGCCTGCCTCAGCCTCCCAAAKTGCTAGGATTATAGGCGTGAGGC 47
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                                                                      NAME/KEY: allele
// LCCATION: 24
// CTHER INFORMATION: 99-21516-293 : polymorphic base G or T
US-09-422-978-2999
                                                                                                                                                                                          Length 47;
                                                                                                                                                                                                                                        6; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION 1435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 9112801.7
FILING DATE: 13-Jun-1991
APPLICATION NUMBER: 9112795.1
FILING DATE: 13-Jun-1991
APPLICATION NUMBER: 9112797.7
FILING DATE: 13-Jun-1991
APPLICATION NUMBER: 9112799.3
FILING DATE: 13-Jun-1991
APPLICATION NUMBER: 05789,067
FILING DATE: 13-Jun-1991
APPLICATION NUMBER: US 07/899,067
FILING DATE: 13-Jun-1991
APPLICATION NUMBER: 1673
REGISTRATION NUMBER: 1673
REGISTRATION NUMBER: 1673
REGISTRATION NUMBER: 96358/PH.36394/US
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
TELEPHONE: (202) 861-3000
TELEPHONE: (202) 861-3000
                                                                                                                                                                                          1.6%; Score 37; DB 1;
85.1%; Pred. No. 24;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 10, Application US/08255889
Patent No. 5525467
GENERAL INFORMATION:
APPLICANT: ANAND, RAKESH
TITLE OF INVENTION: AMPLIFICATION METHODS
NUMBER OF SEQUENCES: 37
CORRESPONDENCE ADDRESS:
ADDRESSEE: CUSHMAN DARBY & CUSHMAN
                                                                                                                                                                                                                                     1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
1.4%; Score 34.2; E
Best Local Similarity 94.3%; Pred. No. 39;
Matches 33; Conservative 2; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: PC-DOS
SOFTWARE: ASCII from WPS-DOS
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/255,889
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: CUSHMAN DARBY & STREET: 1615 L STREET, N.W. CITY: WASHINGTON, D.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Diskette, 3.5'
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (202) 822-0944
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                     Query Match
Best Local Similarity 85.19
Matches 40; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 35
TYPE: Nucleic acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 20036
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                          ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: Linear US-08-255-889-10
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                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 45
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US-09-422-979-2999

Sequence 2999, Application US/09422978

Sequence 2999, Application US/09422978

Sequence 2999, Application US/09422978

Parent No. 6537751

GENERAL INFORMATION:

APPLICANT: Cohen, Daniel

APPLICANT: Chumakov, Ilya

TITLE OF INVENTION: Biallelic markers for use in constructing a high density...

FILE REFERENCE: GENSET. 020CP1

CURRENT FILING DATE: 1999-10-20

EARLIER PILING DATE: 1999-04-21

EARLIER PILING DATE: 1999-04-21

EARLIER PILING DATE: 1999-04-21

EARLIER FILING DATE: 1999-04-21

EARLIER FILING DATE: 1999-04-21

EARLIER FILING DATE: 1998-11-23

EARLIER FILING DATE: 1998-11-23

EARLIER FILING DATE: 1998-04-21
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                                                                 APPLICANT: Larionov, Vladimir L.
APPLICANT: Carinov, Vladimir L.
APPLICANT: Kouprina, Natalay Y.
APPLICANT: Ferkins, Edward L.
TITLE OF INVENTION: TRANSFORMATION-ASSOCIATED RECOMBINATION
TITLE OF INVENTION: CLONING
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSES: Needle & Rosenberg, P.C.
STREET: Suite 1200, 127 Peachtree Street, N.E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 40;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2332 TCGGCCTCCCAAAGTGCTGGGATTACAGGCATGAGCCACC 2371
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        40 TCGGCCTCCCAAAGTGCTGGGATTACAGGCGTGAGCCACC 1
                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                     B: Needle & Rosenberg, P.C.
Suite 1200, 127 Peachtree Street, N.Ε.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
1.6%; Score 38.4; Di
Best Local Similarity 97.5%; Pred. No. 20;
Matches 39; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Patent No. 6391642
GENERAL INPORMATION:
APPLICANT: RESENICK, Michael A.
APPLICANT: Larionov, Vladimir L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 40 bases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQ ID NOS: 11796
SEQ ID NO 2999
                                                                                                                                                                                                                                                                          CITY: Atlanta
CITY: Atlanta
STATE: Georgia
COUNTRY: USA
ZIP: 30303-1811
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy di
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           404-688-9880
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-060-023A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 47
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1.2%; Score 28; DB 1; Length 28;
100.0%; Pred. No. 1e+02;
tive 0; Mismatches 0; Indels
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APPLICANT: Chenchik, Alex
APPLICANT: Jokhadze, George
APPLICANT: Jokhadze, George
APPLICANT: Bibilashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 1375
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
CITY: Menlo Park
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/859,998
FILING DATE: 21-MAY.1997
CLASSIFICATION 1435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
APPLICATION NUMBER:
APPLICATION NUMBER:
APPLICATION NUMBER:
APPLICATION NUMBER:
NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REGISTRATION NUMBER: 37,620
REGISTRATION NUMBER: 09096/002001
TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1.2%; Score 28; DB 1;
100.0%; Pred. No. 1e+02;
cive 0; Mismatches (
                                                                                                                                                                                                                                  ; OTHER INFORMATION: oligonucleotide primer US-08-859-998-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: oligonuclectide primer
                                                                                                                                                                                                                                                                                                                                                                                                  920 GGAGATATGTTGTGAAAGAAGCAGTAGC 947
                                                                                                                                                                                                                                                                                                                                                                                                                                                1 GGAGATATGTTGTGAAGAAGCAGTAGC 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 28, Application US/08859998
Patent No. 5994076
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 415-854-0875
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 28 base pairs
                     TELEFAX: 415-854-0875
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                            Best Local Similarity 100.
Matches 28; Conservative
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Best Local Similarity 100.
Matches 28; Conservative
                                                                                                28 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                 TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM
                                                                                                                                                            TOPOLOGY: linear
MOLECULE TYPE: DNA
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
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    TELEPHONE:
                                                                                                LENGTH:
                                                                                                                                                                                                                                                                                                        Query Match
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US-09-540-699-20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Length 29;
86;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Agrawal, Sudhir
APPLICANT: Agrawal, Sudhir
APPLICANT: Randimalla, Ekambar R.
TITLE OF INVENTION BEGUGO-Cyclic Oligonucleobases
FILE REFERENCE: 99,128-B
CURRENT APPLICATION NUMBER: US/09/540,699
CURRENT APPLICATION NUMBER: US 60/127,138
PRIOR APPLICATION NUMBER: US 60/127,138
PRIOR APPLICATION NUMBER: US 60/174,642
PRIOR PILING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 26
SOFTWARE: PATENTIN UNER: 2000-01-05
SEQ ID NO 20
IENGTH: 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 27, Application US/08859998
Patent No. 5994076
GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
APPLICANT: Jokhadze, George
APPLICANT: Bibilashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 1375
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/859,998
1 GCCTCCCAAAGTGCTGGGATTACAGGYRTGAGCCA 35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1.2%; Score 29; DB
100.0%; Pred. No. 86;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              649 AGGAATCATCGGACTCAGGTACATCTGTG 677
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Fish & Richardson, P.C. STREET: 2200 Sand Hill Road, Suite 100 CITY: Menlo Park STATE: CA COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       29 AGGAATCATCGGACTCAGGTACATCTGTG 1
                                                                                                               Sequence 20, Application US/09540699
Patent No. 6383752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 0909:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE: 21-WAY-1997
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 100.0
Matches 29; Conservative
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US-08-859-998-27
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Gaps

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Length 28;

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TELEFAX: 415-854-0875
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
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    NUMBER OF SEQUENCES: 1375
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single
                                                                                                                                                                    CITY: Menlo Park
STATE: CA
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ò
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                                                                                                                                                                                                                                                                         Sequence 27, Application US/09225928
Patent No. 6352829
GENERAL INFORMATIC Chenchik, Alex
APPLICANT: Chenchik, Alex
Johnadze, George
Bibliashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Bibilashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION: oligonucleotide primer SEQUENCE DESCRIPTION: SEQ ID NO: 27:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/225,928
TILING DATE: 05-Jan-1999
CLASSIFICATION: <unstraint and the companies of the comp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
1.2%; Score 28; DB 1;
Best Local Similarity 100.0%; Pred. No. 1e+02;
Matches 28; Conservative 0; Mismatches 0
1204 CCTTAGCTGACTATTGGAAATGCACTTC 1231
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   920 GGAGATATGTTGTGAAAGAAGCAGTAGC 947
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 GGAGATATGTTGTGAAAGAAGCAGTAGC 28
                                                                                28 cérradergacrarragaaargeacrie 1
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Sequence 28, Application US/09225928
Patent No. 6352829
GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
JOKHAGZE, George
Bibilashvilli, Robert
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: 415-322-5070
TELEFAX: 415-854-0875
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 1375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 94025
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 50
US-09-225-928-28/c
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ORGANIEM: Artificial Sequence
PEATURE:
OTHER INFORMATION: Description of Artificial Sequence:Oligonucleotide
OTHER INFORMATION: that is complementary to a portion of the human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ;
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                                                                                                                 COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FASTENCY for Windows Version 2.0
CURRIN APPLICATION DATA:
APPLICATION NUMBER: US/09/225,928
FILING DATE: 05-Jan-1999
CLASSIFTCATION: «Unknown>
PRIOR APPLICATION CURREN: 08/859,998
FILING DATE: 21-MAY-1997
ATTORNEY/AGRAT INFORMATION:
NAMME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 37,620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-16; Sequence 16, Application US/09540699; Sequence 16, Application US/09540699; Patent No. 6383752; GENERAL INFORMATION; Patent No. 6383752; GENERAL INFORMATION; Patent No. 6383762; TILLE OF INVENTION; Pacudo-Cyclic Oligonucleobases; FILE REFRENCE: 99,128-B; CURRENT APPLICATION NUMBER: US/09/540,699; CURRENT PILING DATE: 2000-03-31; PRIOR APPLICATION NUMBER: US 60/174,642; PRIOR FILING DATE: 2000-01-05; NUMBER OF SEQ ID NOS: 26; SOFTWARE: PatentIn Ver: 2.0; LENGTH:: 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; SEQUENCE DESCRIPTION: Oligonucleotide primer; SEQUENCE DESCRIPTION: SEQ ID NO: 28:
ADDRESSEE: Fish & Richardson, P.C. STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
. 1e+02;
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1.2%; Score 28; DB
Best Local Similarity 100.0%; Pred. No. 1e+
Matches 28; Conservative 0; Mismatches
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TELEPHONE: 415-322-5070
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CITY: Menlo Park
                                                                Jokhadze,
                                                                                                                                                                                                                                                                                         COUNTRY: US
Patent No. 6489455
GENERAL INFORMATION:
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US-09-304-232-196
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100.0%; Pred. No. 1e+02;
tive 0; Mismatches 0; Indels
                                                             Query Match 1.2%; Score 28; DB 1; Length 28; Best Local Similarity 100.0%; Pred. No. 18+02; Matches 28; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                              JOKNBALL, ROBERT BIDLIASHVILLI, ROBERT TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT INDEACULE
COMPUTER: IBM Compatible
COPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/225,201B
FILING DATE: 05-Jan-1999
CLASSIFICATION DATA:
APPLICATION DATA:
AREFERENCE DATA:
NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 09096/002001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; SEQUENCE DESCRIPTION: Oligonucleotide primer; SEQUENCE DESCRIPTION: SEQ ID NO: 27: US-09-225-201B-27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                    671 ATCTGTGAGTGAGAACAGGTGTCACCTT 698
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                920 GGAGATATGTTGTGAAAGAAGCAGTAGC 947
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                                                                                                                                                                                                                                                                                                                                                                         Chenchia, ...
Jokhadze, George
                                                                                                                                                                                                                                                                                                       Sequence 27, Application US/09225201B Patent No. 6489455 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 28, Application US/09225201B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 415-854-0875 INFORMATION FOR SEQ ID NO: 27:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Chenchik, Alex
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE CHARACTERISTICS
OTHER INFORMATION: MDM2 mRNA
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Best Local Similarity 100.
Matches 28; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: DNA FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 53
US-09-225-201B-28/c
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US-09-225-201B-27
    ; US-09-540-699-16
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Sequence 196, Application US/09304232

Sequence 196, Application US/09304232

GENERAL INFORMATION:
APPLICANT: Fan, Jian Bing
APPLICANT: Case Western Kenneth
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Hypertension
FILE REFERENCE: 018847-01421005
CURRENT APPLICATION NUMBER: US/09/304,232
CURRENT APPLICATION NUMBER: US/09/304,232

CURRENT PILING DATE: 1999-05-03

EARLIER PILING DATE: 1999-05-07

NUMBER OF SEQ ID NOS: 909
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                                             Bibilashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ) OTHER INFORMATION: oligonucleotide primer ; SEQUENCE DESCRIPTION: SEQ ID NO: 28: US-09-225-201B-28
                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM COMPATIBLE
COMPUTER: IBM COMPATIBLE
COMPUTER: IBM COMPATIBLE
SOFTWARE: PESESEG for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/225,201B
FILLING DATE: 05-Jan-1999
CLASSIFICATION: «Unknown»
PRIOR APPLICATION DATA:
                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               09096/002001
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Best Local Similarity 100.0%; Pred. No. 157
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/08/859,998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1204 CCTTAGCTGACTATTGGAAATGCACTTC 1231
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 09
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE: 21-MAY-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                          ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 415-854-0875
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 28 base pairs
APPLICANT: Chenchik, Alex
                                                                                                                               NUMBER OF SEQUENCES: 1375
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
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; SEQ ID NO 571

LENGTH: 29

TYPE: DNA

ORGANISM: Artificial Sequence
FRATURE:

COTHER INFORMATION: HSTSCGENE 3838
US-09-304-232-571
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US-09-540-699-18
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Sequence 513, Application US/09304232

Ratent No. 6525185

GENERAL INFORMATION:

APPLICANT: Fan, June Bing

APPLICANT: Case Western Kenneth

APPLICANT: Case Western Keserve University School of Medicine

APPLICANT: Affwerix, Inc.

TITLE OF INVENTION: Polymorphisms Associated With

TITLE OF INVENTION: Hypercension

FILE REFERENCE: 01854-034210US

CURRENY FILING DATE: 1999-05-03

CURRENY FILING DATE: 1999-05-03

EARLIER FILING DATE: 1998-05-07

NUMBER OF SEQ ID NOS: 909

SEQ ID NO 513

EARLIER PAILSON WHOMER: US 60/084,641

SOFTHARE: PSSEC for Windows Version 3.0

SEQ ID NO 513

EMENTER: PSSEC FOR WINDOWS OF SECOND OF OF SECON
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APPLICANT: Far. Jan Bing
APPLICANT: Chakravarti, Aravinda
APPLICANT: Halushka, Marc Kenneth
APPLICANT: Halushka, Marc Kenneth
APPLICANT: Halushka, Marc Kenneth
APPLICANT: Affwertix, Inc.
ITLE OF INVENTION: Polymorphisms Associated With
ITLE OF INVENTION: Polymorphisms Associated With
ITLE OF INVENTION: Polymorphisms Associated With
ITLE OF INVENTION: 1994-1003
FILE REPRENCE: 11994-05-03
CURRENT APPLICATION NUMBER: US 60/084,641
EARLIER FILING DATE: 1998-05-07
NUMBER OF SEQ ID NOS: 909
SOFTWARE: FastSEQ for Windows Version 3.0
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1.1%; Score 27; DB 1; Length 29;
Best Local Similarity 93.1%; Pred. No. 1.1e+02;
Matches 27; Conservative 1; Mismatches 1; Indels
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1.1%; Score 27; DB 1; I
Best Local Similarity 93.1%; Pred. No. 1.1e+02;
Matches 27; Conservative 1; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2332 TCGGCCTCCCAAAGTGCTGGGATTACAGG 2360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2281 ACCGTGTTAGCCAGGATGGTCTCGATCTC 2309
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SOFTWARE: FastSEQ for Windows Version 3.0
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Patent No. 6525185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ) OTHER INFORMATION: GLUT4EX11 941 US-09-304-232-513
                                                                                                                 TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                  ; OTHER INFORMATION: APOC4 1287
US-09-304-232-196
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Artificial Sequence
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                                    SEQ·ID NO 196
LENGTH: 29
                                                                                                                                                                                                  FEATURE:
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                                                                                                            Gaps
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     Length 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 270, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
ITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIF: ".....
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
Query Match
1.1%; Score 27; DB 1; 1
Best Local Similarity 93.1%; Pred. No. 1.1e+02;
Matches 27; Conservative 1; Mismatches 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street CITY: Marlton STATE: NJ STATE: NJ ZIP: 08053
                                                                                                                                                                                                   2339 CCCAAAGIGCIGGGATIACAGGCAIGAGC 2367
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CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
ATTORNEY AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 12,257
REFERENCE/DOCKET NUMBER: 12FP+0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION:
TELECOMMUNI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   415 TGAAGTTATTAAAGTCTGTTGGTGCA 440
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 26 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
ANTI-SENSE: Yes
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APPLICANT: Fan, Jian Bing
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US-09-304-232-156
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GENERAL INFORMATION:

APPLICANT: Fan, Jian Bing

APPLICANT: Chakravarti, Aravinda

APPLICANT: Halushka, Marc Kenneth

APPLICANT: Gase Western Reserve University School of Medicine

APPLICANT: Affymetrix, Inc.

TITLE OF INVENTION: POlymorphisms Associated With

TITLE OF INVENTION: Hypertension

FILE REPERENCE: 018647-034210US

CURRENT FILING DATE: 1999-05-03

CURRENT APPLICATION NUMBER: US/09/304,532

CURRENT FILING DATE: 1999-05-03

EARLIER FILING DATE: 1999-05-03

SOFTWARE: FastSEQ ID NOS: 909

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 503
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: MDM2 mRNA OTHER INFORMATION: forward primer; +311 to +366).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 29;
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                                                                  APPLICANT: Agrawal, Sudhir
APPLICANT: Kandimalla, Ekambar R.
TITLE OF INVENTION: Peeudo-Cyclic Oligonucleobases
FILE REFERENCE: 99,128-B
CURRENT APPLICATION NUMBER: US/09/540,699
CURRENT FILING DATE: 2000-03-31
PRIOR PILING DATE: 1999-03-31
PRIOR APPLICATION NUMBER: US 60/127,138
PRIOR FILING DATE: 1999-01-31
PRIOR APPLICATION NUMBER: US 60/174,642
PRIOR FILING DATE: 2000-01-05
NUMBER OF SEQ. ID NOS: 26
SOFTWARE: PatentIN Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2345 GTGCTGGGATTACAGGCATGAGCCACCG 2372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 GTGCTGGGATTACARGCGTGAGCCACCG 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        622 ACAGGAACTTGGTAGTAGTCAATCAG 647
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 ACAGGAACTTGGTAGTGATCAG 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 503, Application US/09304232 Patent No. 6525185
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Patent No. 6407062
Sequence 18, Application US/09540699
Patent No. 6383752
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: GLUT4EX11 1005
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-304-232-503
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                                                                                                                                                                                                                                                                                                                                                                                SEQ ID NO 18
LENGTH: 26
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APPLICANT: Sherr, Charles J
APPLICANT: Sherr, Charles J
APPLICANT: Guelle, Dawn E
APPLICANT: Woule, Jawn E
APPLICANT: Rouses, Martine F.
APPLICANT: Rouses, Martine F.
APPLICANT: Frederique, Zindy
TITLE OF INVENTION: AAF-19, A NOVEL REGULATOR OF THE MAMALIAN CELL CYCLE
FILE REFERENCE: 1340-1-023 CIP 1
CURRENT APPLICATION NUMBER: US/09/480,718
CURRENT FILING DATE: 2000-01-07
EARLIER FILING DATE: 1998-08-06
NUMBER OF SEQ ID NOS: 48
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 35
LENGTH: 30
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| Patent No. 6252185 |
| GENERAL INFORMATION |
| APPLICANT: Fan, Jian Bing |
| APPLICANT: Chakravarti, Aravinda |
| APPLICANT: Chakravarti, Aravinda |
| APPLICANT: Case Western Reserve University School of Medicine |
| APPLICANT: Case Western Reserve University School of Medicine |
| APPLICANT: APPLICANT: APPLICANT: Inc. |
| TITLE OF INVENTION: Polymorphisms Associated With |
| TITLE OF INVENTION: Hypertension |
| TITLE REFERENCE: 010547-03421005 |
| CURRENT APPLICATION NUMBER: US/09/304,232 |
| CURRENT PILING DATE: 1999-05-03 |
| BARLIER APPLICATION NUMBER: US 60/084,641 |
| EARLIER FILING DATE: 1999-05-07 |
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 30;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 25.8; DB 1;
Pred. No. 1.3e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1.1%; Scor.
93.1%; Pred. No. 1...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                312 AIGIGCAAIACCAACAIGICTGIACCIAC 340
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SOFTWARES FastSEQ for Windows Version 3.0
SEQ ID NO 156
LENGTH: 29
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: APOCIEX1 1020 US-09-304-232-156
                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Artificial Sequence
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Matches 27; Conservative
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LENGTH: 25 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOCY: Linear
ANTI-SENSE: Yes
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OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
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US-09-280-805-271
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US-09-280-805-271
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CITY: Ma
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APPLICANT: Fan, Jian Bing
APPLICANT: Chakravarti, Aravinda
APPLICANT: Chakravarti, Aravinda
APPLICANT: Case Western Reserve University School of Medicine
APPLICANT: AFFMerrix, Inc.
ITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: 1999-003
FILE REFERENCE: 018547-034210US
CURRENT APPLICATION NUMBER: US,09/304,232
CURRENT FILING DATE: 1999-06-03
EARLIER APPLICATION NUMBER: US 60/084,641
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: Chakravarti, Aravinda
: Halushka, Marc Kenneth
: Case Western Reserve University School of Medicine
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1.1%; Score 25.4; DB 1; Length 29;
Best Local Similarity 89.7%; Pred. No. 1.4e+02;
Matches 26; Conservative 1; Mismatches 2; Indels
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                      APPLICANT: Halushka, Marc Kenneth
APPLICANT: Case Western Reserve University School of
APPLICANT: Case Western Reserve University School of
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Hypertension
FILE REPRENCE: 0.18547-0.3421003
CURRENT PELING DATE: 1999-05-03
EARLIER APPLICATION NUMBER: US 60/084,641
EARLIER PILING DATE: 1998-05-07
NUMBER OF SEQ ID NOS: 909
NUMBER OF SEQ ID NOS: 909
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2326 CCCACCTCGGCCTCCCAAAGTGCTGGGAT 2354
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 512
LENGTH: 29
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US-09-304-232-705
; Sequence 705. Application US/09304232
; Patent No. 6525185
; GENERAL INFORMATION:
; APPLICANT: Fan, Jian Bing
; APPLICANT: Chakravarti, Aravinda
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-304-232-512
; Sequence 512, Application US/09304232
; Patent No. 6525185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: GLUT4EX11 935
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CTHER INFORMATION: APOC4 1281
US-09-304-232-195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                            ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-304-232-512
                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
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APPLICANT: Halushka, Marc Kenneth
APPLICANT: Case Western Reserve University School of Medicine
APPLICANT: Case Western Reserve University School of Medicine
TITLE OF INVENTION: POLYMORPHISMS
TITLE OF INVENTION: Hypertension
FILE REFERENCE: 018547-03410US
CURRENT APPLICATION NUMBER: US/09/304,232
CURRENT APPLICATION NUMBER: US 60/084,641
EARLIER FILING DATE: 1998-05-07
NUMBER OF SEQ ID NOS: 909
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 705
LENGTH: 29
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Patent No. 6184212
GENERAL INFORMATION
GENERAL INFORMATION
APPLICANT: Graham, Brett P. Monia
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DIEKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.1%; Score 25.4; DB 1;
89.7%; Pred. No. 1.4e+02;
tive 1; Mismatches 2;
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
PRPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                              ) OTHER INFORMATION: PGISEX10 3186
US-09-304-232-705
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISE
TELECOMMUNICATION
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 89./*
The 26, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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Gaps
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Sequence 193, Application US/09304232

Sequence INFORMATION:
GENERAL INFORMATION:
APPLICANT: Fan, Julia Bing
APPLICANT: Case Western Keserve University School of Medicine
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Hypertenaion
TITLE OF INVENTION: Hypertenaion
TITLE MAPLICATION NUMBER: US/09/304,232
CURRENT APPLICATION NUMBER: US/09/304,232
CURRENT APPLICATION NUMBER: US 60/084,641

EARLIER FILING DATE: 1999-05-07

NUMBER OF SEQ ID NOS: 909
SEQ ID NO 193
LENGTH: 29
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| Batent No. 6225185 |
| GENERAL INFORMATION |
| APPLICANT: Ran, Jian Bing |
| APPLICANT: Halushka, Marc Kenneth |
| APPLICANT: Case Western Reserve University School of Medicine |
| APPLICANT: Case Western Reserve University School of Medicine |
| APPLICANT: Affymetrix, Inc. |
| TITLE OF INVENTION: Polymorphisms Associated With |
| TITLE OF INVENTION: Hypertension |
| TITLE OF INVENTION: Hypertension |
| TITLE OF INVENTION UNMERR: US/09/304,232 |
| CURRENT APPLICATION NUMBER: US 60/084,641 |
| EARLIER FILING DATE: 1998-05-07 |
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                              1; Indels
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89.3%; Pred. No. 1.6e+02;
     92.6%; Pred. No. 1.5e+02;
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                                 1; Mismatches
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                                                                                    2274 GGGTTTCACCGTGTTAGCCAGGATGGT 2300
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SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 698
LENGTH: 29
                                                                                                                3 GGATTTCACCGTRTTAGCCAGGATGGT 29
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ORGANISM: Artificial Sequence
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OTHER INFORMATION: APOC4 1150
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                              25; Conservative
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Best Local Similarity
Best Local Similarity
Matches 25; Conserv
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US-09-304-232-193
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US-09-304-232-698
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Batent No. 6525185

GENERAL INFORMATION:

APPLICANT: Fan, Jian Bing
APPLICANT: Halushka, Marc Kenneth
APPLICANT: Case Western Reserve University School of Medicine
APPLICANT: Afymetrix, Inc.
TITLE OF INVENTION: POlymorphisms Associated With
TITLE OF INVENTION: Hypertension
FILE REFERENCE: 01854-034210US
CURRENT FILING DATE: 1999-05-03

CURRENT FILING DATE: 1999-05-03

BARLIER FILING DATE: 1999-05-07

NUMBER OF SEQ ID NOS: 909

SOFTWARE: FastSEQ for Windows Version 3.0

SEQ ID NO 703

LINGTH: 29
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                                                                                                                                                                                                                                                                                      Sequence 4, Application US/09837149

| Sequence 4, Application US/09837149
| Patent No. 6448014
| Patent No. 6448014
| APPLICANT: Cloyd, Miles W.
| APPLICANT: Cloyd, Miles W.
| APPLICANT: Chen Application Assays Specific for TITLE OF INVENTION: Integrated Retroviruses
| TITLE OF INVENTION: Integrated Retroviruses
| FILE REPRENCE: D6285
| CURRENT APPLICATION NUMBER: US/09/837,149
| PRIOR APPLICATION NUMBER: US 60/198,884
| PRIOR FILING DATE: 2000-04-18
| NUMBER OF SEO.ID NOS: 4
| SEQ.ID NO 4
| SEQ.ID NOS: 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: primer
OTHER INFORMATION: primer for the Alu sequence in the human
OTHER INFORMATION: chromosomal DNA
                            Length 25;
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                            DB 1; Le
1.6e+02;
                      Query Match 1.1%; Score 25; DB Best Local Similarity 100.0%; Pred. No. 1.6 Matches 25; Conservative 0; Mismatches
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                                                                                                                                    355 CCACCTCACAGATTCCAGCTTCGGA 379
                                                                                                                                                                                    1 ccaccicacagariccagcricgga 25
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US-09-304-232-703
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JUREAL AND CANALION:

APPLICANT: Sherr, Charles J
APPLICANT: Quelle, Dawn E
APPLICANT: Quelle, Dawn E
APPLICANT: Quelle, Dawn E
APPLICANT: Quelle, Zindy
APPLICANT: Roussel, Martine F.
APPLICANT: Rederique, Zindy
TITLE OF INVENTION: ARF-19, A NOVEL REGULATOR OF THE MAMMALIAN CELL CYCLE
TITLE OF INVENTION: ARF-19, A NOVEL REGULATOR OF THE MAMMALIAN CELL CYCLE
CURRENT APPLICATION NUMBER: US/09/480,718
CURRENT APPLICATION NUMBER: US/09/855
EARLIER APPLICATION NUMBER: 09/129,855
EARLIER PILING DATE: 1998-08-06
NUMBER OF SEQ ID NOS: 48
SEQ ID NO 28
LENGTH: 30
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:

APPLICANT: Fan, Jian Bing
APPLICANT: Chakravarti, Aravinda
APPLICANT: Halushka, Marc Kenneth
APPLICANT: Halushka, Marc Kenneth
APPLICANT: Affwetrix, Inc.
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Hypertension
FILE REFERENCE: 018547-03421005
CURRENT PILING DATE: 1999-05-03
CURRENT FILING DATE: 1998-05-03
EARLIER FILING DATE: 1998-05-07
NUMBER OF SEQ ID NOS: 909
SOFTWARE: FastSEQ for Windows Version 3.0
LENGTH: 29
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1.0%; Score 24.2; DB 1;
Best Local Similarity 89.7%; Pred. No. 1.7e+02;
Matches 26; Conservative 0; Mismatches 3;
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Pred. No. 1.7e+02;
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; Sequence 706, Application US/09304232
; Patent No. 6525185
                      Sequence 28, Application US/09480718
Patent No. 6407062
GENERAL INFORMATION:
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US-09-304-232-79/c
; Sequence 79, Application US/09304232
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US-09-304-232-706
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ORGANISM: Artificial Sequence
FEATURE:
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Best Local Similarity 92.3%;
Matches 24; Conservative
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APPLICANT: DePinho, Robert A.
TITLE OF INVENTION: A METHOD OF INHIBITING CELL PROLIFERATION USING AN ANTI-ONCOGENE
FILE REFERENCE: 96700/469
CURRENT APPLICATION NUMBER: US/09/044,602
CURRENT FILING DATE: 1998-03-19
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin version 3.1
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GENERAL INFORMATION:

APPLICANT: Chakravarti, Aravinda

APPLICANT: Chakravarti, Aravinda

APPLICANT: Halushka, Marc Kenneth

APPLICANT: Affymetrix, Inc.

TITLE OF INVENTION: Polymorphisms Associated With

TITLE OF INVENTION: Hypertension

FILE REPERENCE: 018647-03421008

CURRENT APPLICATION NUMBER: US/09/304,232

CURRENT PILING DATE: 1999-05-03

EARLIER APPLICATION NUMBER: US 60/084,641

EARLIER PILING DATE: 1998-05-07

NUMBER OF SEQ ID NOS: 909

SOFTWARE: FastSEQ for Windows Version 3.0

LENGTH: 29
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1.0%; Score 24.2; DB 1; Length 29;
Best Local Similarity 89.7%; Pred. No. 1.7e+02;
Matches 26; Conservative 0; Mismatches 3; Indels
  Indels
  2;
  1; Mismatches
                                               2140 GGGTGATCTTGGCTCACTGCAAGCTCTG 2167
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CTHER INFORMATION: primer for MDM2 mutant US-09-044-602-1
                                                                                                                                                                                                         Sequence 704, Application US/09304232
Patent No. 6525185
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Patent No. 6613750
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CTHER INFORMATION: PGISEXIO 3140

US-09-304-232-704
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25; Conservative
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LENGTH: 29
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US-09-304-232-589/c
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LENGTH: 29
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APPLICANT: Fan, Jian Bing
APPLICANT: Chakravarti, Aravinda
APPLICANT: Haluebhka, Marc Kenneth
APPLICANT: Haluebhka, Marc Kenneth
APPLICANT: Gae Western Reserve University School of Medicine
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Hypertension
FILE REFERENCE: 018647-034210US
CURRENT FILING DATE: 1999-05-03
CURRENT FILING DATE: 1999-05-03
EARLIER FILING DATE: 1998-05-07
NUMBER OF SEQ ID NOS: 909
SOFTWARE: PastSEQ for Windows Version 3.0
SEQ ID NO 158
                                      APPLICANT: CLASS AND AND AND AND APPLICANT: Class Western Reserve University School of Medicine APPLICANT: Case Western Reserve University School of Medicine APPLICANT: Affectix, Inc.
ITILE OF INVENTION: Polymorphisms Associated With TITLE OF INVENTION: Hypertension FILE REFERENCE: 0.1847-0421008
CURRENT APPLICATION NUMBER: US 60/084,641
EARLIER APPLICATION NUMBER: US 60/084,641
EARLIER PILING DATE: 1999-05-07
NUMBER OF SEQ ID NOS: 909
SOFTWARE: FastSEQ for Windows Version 3.0
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1.0%; Score 23.8; DB 1;
Best Local Similarity 86.2%; Pred. No. 1.8e+02;
Matches 25; Conservative 1; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 23.8; DB 1;
Pred. No. 1.8e+02;
1; Mismatches 3;
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Patent No. 6525185
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Patent No. 6525185
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US-09-304-232-158
                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: AE1EX20 1679
APPLICANT: Fan, Jian Bing
APPLICANT: Chakravarti, Aravinda
                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 1.0%;
Best Local Similarity 86.2%;
Matches 25; Conservative
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LENGTH: 29
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APPLICANT: Chakravarti, Aravinda
APPLICANT: Chakravarti, Aravinda
APPLICANT: Chakravarti, Aravinda
APPLICANT: Halushka, Marc Kenneth
APPLICANT: Affmetix, Inc.
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Polymorphisms Associated With
TITLE REFRENCE: 018547-03421005
CURRENT FILING DATE: 1999-05-03
EARLIER APPLICATION NUMBER: US 60/084,641
EARLIER APPLICATION NUMBER: US 60/084,641
NUMBER OF SEQ ID NOS: 909
SOFTWARE: PREUSEQ for Windows Version 3.0
APPLICANT: Fan, Jian Bing
APPLICANT: Chakravarti, Aravinda
APPLICANT: Chakravarti, Aravinda
APPLICANT: Halushka, Marc Kenneth
APPLICANT: Halushka, Marc Kenneth
APPLICANT: Affwetrix, Inc.
TILLE OF INVENTION: Polymorphisms Associated With
TILLE OF INVENTION: Polymorphisms Associated With
TILLE OF INVENTION: Pypertension
FILE REPREBRUCE: 015547-034210US
CURRENT APPLICATION NUMBER: US 60/084,641
EARLIER PILING DATE: 1999-05-03
EARLIER PILING DATE: 1998-05-07
NUMBER OF SEQ ID NOTE: 1998-05-07
SOFWARE: FastSEQ for Windows Version 3.0
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Pred. No. 1.8e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 23.8; DB 1;
Pred. No. 1.8e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1; Mismatches
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; Sequence 514, Application US/09304232
Patent No. 6525185
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: GLUT4EX11 963
US-09-304-232-514
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86.2%;
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US-09-304-232-185
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Best Local Similarity 86.2%;
Matches 25; Conservative
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ORGANISM: Artificial Sequence
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Best Local Similarity 86.2
Matches 25; Conservative
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TYPE: DNA

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Kouprina, Natalay Y.
Perkins, Edward L.
VENTION: TRANSFORMATION-ASSOCIATED RECOMBINATION
PENTION: CLONING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 23.6; DB 1; Length 30;
Pred. No. 1.8e+02;
0; Mismatches 4; Indels
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; Sequence 4, Application US/09385917
; Patent No. 6294546
; Patent No. 6294546
; Patent No. 6294546
; APPLICANT: Rosen, Glenn
; APPLICANT: Rosen, Glenn
; TITLE OF INVENTION: USES OF DITERPENOID TRIEPOXIDES AS AN TITLE OF INVENTION: ANTI-PROLIFERATIVE AGENT
; TITLE OF INVENTION: ANTI-PROLIFERATIVE AGENT
; CURRENT APPLICATION NUMBER: US/09/385,917
; CURRENT PILLING DATE: 1999-08-30
; NUMBER OF SEQ ID NOS: 8
; SOFTWARE: PastSEQ for Windows Version 3.0
; SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/060,023A
FILING DATE: April 14, 1998
PRIOR APPLICATION NUMBER: PCT/US96/11478
FILING DATE: UDLY 9, 1996
ATTORNEY/AGENT INFORMATION:
                                                                               TITLE OF INVENTION: TRANSFORMATION-ASSOCIATED RECOM-
TITLE OF INVENTION: CLONING
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSE: Needle & Rosenberg, P.C.
STREET: Suite 1200, 127 Peachtree Street, N.E.
CITY: Atlanta
STATE: Georgia
COUNTRY: USA
ZIP: 30303-1811
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
1.0%; Score 23; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 2.1e+02;
Matches 23; Conservative 0; Mismatches 0;
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             Larionov, Vladimir L.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Perryman, David G.
REGISTRATION NUMBER: 33,438
REFRENCE/DOCKET NUMBER: 14014
TELECOMMUNICATION INFORMATION:
TELEPHONE: 404-688-070
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
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Best Local Similarity 86.7%;
Matches 26; Conservative
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STRANDEDNESS: single
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CORGANISM: H. sapiens
US-09-385-917-4
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MOLECULE TYPE:
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                                        APPLICANT:
APPLICANT:
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APPLICANT: Fan, Jian Bing
APPLICANT: Chakravarti, Aravinda
APPLICANT: Halubhka, Marc Kenneth
APPLICANT: Halubhka, Marc Kenneth
APPLICANT: Affwmetrix, Inc.
ITLE OF INVENTION: Hyptension
ITLE OF INVENTION: Hyptension
FILE REFERENCE: 018547-034210US
CURRENT APPLICATION NUMBER: US/09/304,232
CURRENT APPLICATION NUMBER: US/09/304,641
EARLIER APPLICATION NUMBER: US 60/084,641
BARLIER PILING DATE: 1998-05-07
NUMBER OF SEQ ID NOS: 909
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO S89
LENGTH: 29
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APPLICANT: Chakravarti, Aravinda
APPLICANT: Chakravarti, Aravinda
APPLICANT: Halushka, Marc Kenneth
APPLICANT: Case Western Reserve University School of Medicine
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: 1919-05-03
CURRENT APPLICATION NUMBER: US/09/304,232
CURRENT FILING DATE: 1998-05-07
BARLIER FILING DATE: 1998-05-07
NUMBER OF SEQ ID NOS: 909
SOFTWARE: FRAESEQ for Windows Version 3.0
SOFTWARE: PRAESEQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1.0%; Score 23.8; DB 1;
86.2%; Pred. No. 1.8e+02;
iive 1; Mismatches 3;
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Patent No. 6525185
GENERAL INFORMATION:
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Patent No. 6391642
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: PGISEX10 1500 US-09-304-232-685
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Resnick, Michael A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; FEATURE:
; OTHER INFORMATION: IAPPEX3 848
US-09-304-232-589
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ORGANISM: Artificial Sequence
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Best Local Similarity 86.2%
Matches 25; Conservative
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US-09-304-232-685/c
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US-09-060-023A-4
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FEATURE:

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Length 23; 0; Indels

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Best Local Similarity
Matches 24; Conserva
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    SENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                Sequence 4, Application US/09884898
Patent No. 6537984
GENERAL INFORMATION:
APPLICANT: Rosen, Glenn D.
APPLICANT: Lennox, Edwin S.
APPLICANT: Lennox, Edwin S.
TITLE OF INVENTION: USES OF DITERPENOID TRIEPOXIDES AS AN TITLE OF INVENTION: WHERE TO STANDSOIN
FILE REFERENCE: STANDSOIN
CURRENT APPLICATION NUMBER: US/09/884,898
CURRENT APPLICATION NUMBER: US/09/884,898
CURRENT FILING DATE: 2001-06-19
PRIOR PELICATION NUMBER: 09/385,917
PRIOR FILING DATE: 1005: 8
NUMBER OF SEQ IN NOS: 8
SOFTWARE: FastSEQ for Windows Version 4.0
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Patent No. 6599499

GENERAL INFORMATION:
APPLICANT: Rosen, Glenn D.
APPLICANT: Lennox, Edwin S.
APPLICANT: Lennox, Edwin S.
TITLE OF INVENTION: Uses of Diterpenoid Triepoxides as an TITLE OF INVENTION: Anti-Proliferative Agent; FILE REFERENCE: STAN096CIP; CURRENT APPLICATION NUMBER: US/09/935,794

CURRENT APPLICATION NUMBER: 09/385,917

PRIOR FILING DATE: 2001-08-22

PRIOR FILING DATE: 1999-08-30

NUMBER OF SEQ IN NOS: 16

SOFTWARE: PastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 1.0%; Score 23; DB 1; Length 23; Best Local Similarity 100.0%; Pred. No. 2.1e+02; Matches 23; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1.0%; Score 23; DB 1; Length 23;
100.0%; Pred. No. 2.1e+02;
tive 0; Mismatches 0; Indels
1374 GAGGCTTTGATGTTCCTGATTG 1396
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; Sequence 66, Application US/08859998
; Patent No. 5994076
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.
Matches 23; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TYPE: DNA
; ORGANISM: H. sapiens
US-09-884-898-4
                                                                                                                                  US-09-884-898-4/C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-935-794-4/c
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LENGTH: 23
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LENGTH: 23
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##PRICONT: Chemistry
##PRICONT: Chemistry
##PRED: ADDRESSER: Fish & Rechardson, P.C.
##PRED: ADDRESSER: Fish & R
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MEDIUM TYPE: Diskette
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INFORMATION FOR SEQ ID NO: 66:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              LENGTH: 26 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 1.0%;
Best Local Similarity 92.3%;
Matches 24; Conservative
                                                                                                                  Query Match
Best Local Similarity 92.3%;
Matches 24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: US
ZIP: 94025
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES:
    TYPE: nucleic acid
STRANDEDNESS: single
                                                   linear
                                                   ,
US-09-367-750-10
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Patent No. 643639

GENERAL INFORMATION:

APPLICANT: Kiefer, Michael C.

APPLICANT: Ossina, Natalya K.

TITLE OF INVENTION: Bak PROMOTER EXPRESSION SYSTEM

NUMBER OF SEQUENCES: 14

CORRESPONDENCE ADDRESS:

ADDRESSEE: LXR BIOTECHNOLOGY INC.

STREET: 3095 Richmond Parkway, Suite 213

CCUNTRY: Richmond

STATE: CAR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: PETENT REDS/HS DOS
SOFTWARE: PETENT RELEAGE #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/367,750
FILING DATE: 0.05C-1999
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: BROWN, THEFERA A.
REGISTRATION NUMBER: 32,547
REFERENCE/DOCKET NUMBER: 4147-14-PUS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; SEQUENCE DESCRIPTION: Oligonucleotide primer ; SEQUENCE DESCRIPTION: SEQ ID NO: 66: US-09-225-928-66
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FascESD for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/225,928
FILING DATE: 05-Jan-1999
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/859,998
FILING DATE: 21-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 09096/002001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2342 AAAGTGCTGGGATTACAGGCATGAGC 2367
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COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
CMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 26 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 415-854-0875
INFORMATION FOR SEQ ID NO: 66:
                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-367-750-10
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                                                IndelB
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                                                                                                                                                                                                                                                                                                            APPLICANT: Chenchik, Alex
Jokhadze, George
Bibliashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; CTHER INFORMATION: Oligonucleotide primer; SEQUENCE DESCRIPTION: SEQ ID NO: 66: US-09-225-201B-66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/225,201B
FILING DATE: 05-Jan-1999
CLASSIFICATION: cUnknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/859,998
FILING DATE: 21-MAY-1997
ATTORNEY/AGRYI INFORMATION:
NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 09096/002001
TELECOMMUNICATION: 1NFORMATION:
TELECOMMUNICATION: 415-322-5070
Score 22.8; DB 1;
Pred. No. 2.1e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 22.8; DB 1;
Pred. No. 2.1e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
CITY: Menlo Park
STATE: CA
                                                                                         2264 AGTAGAGACAGGGTTTCACCGTGTTA 2289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2342 AAAGTGCTGGGATTACAGGCATGAGC 2367
                                                                                                                                  1 AGTAGAGACGGGTTTCACCATGTTA 26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26
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                                                                                                                                                                                                                          US-09-225-201B-66; Sequence 66, Application US/09225201B Fatent No. 6489455; GENERAL INFORMATION:
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Sequence 6, Application US/08454557C
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US-08-340-426D-6/c
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                        Sequence 511, Application US/09304232

Fatent No. 6525185

GENERAL INFORMATION:

APPLICANT: Fan, Jian Bing

APPLICANT: Cask wareti, Aravinda

APPLICANT: Gase Western Kenneth

APPLICANT: Affymetrix, Inc.

TITLE OF INVENTION: Polymorphisms Associated With

TITLE OF INVENTION: Polymorphisms Associated With

TITLE OF INVENTION: Hypertension

FILE REFERENCE: 018547-03421008

CURRENT APPLICATION NUMBER: US 60/084,641

EARLIER FILING DATE: 1999-05-03

KARLIER PELLING DATE: 1999-05-07

NUMBER OF SEQ ID NOS: 909

SOFTWARE FEALSE FREESE F
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Patent NO. 6525185

GENERAL INFORMATION:

APPLICANT: Fan, Jian Bing

APPLICANT: Halushka, Marc Kenneth

APPLICANT: Affymetrix, Inc.

TILE OF INVENTION: Polymorphisms Associated With

TILE REFERENCE: 018547-034210US

CURRENT FILING DATE: 1999-05-03

BARLIER APPLICATION NUMBER: US 60/084,641

FARLIER APPLICATION NUMBER: US 60/084,641

BARLIER APPLICATION NUMBER: 999-05-03

BARLIER PAPEL 90 DATE: 1999-05-07

NUMBER OF EQ ID NOS: 909

SOFTWARE: FastSEQ for Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 1.0%; Score 22.8; DB 1; Length 29; Best Local Similarity 85.7%; Pred. No. 2e+02; Matches 24; Conservative 1; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2271 ACAGGGTTTCACCGTGTTAGCCAGGATG 2298
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2106 GAGTCTTGCTCTGTTACCCAGGCTGGAG 2133
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      , OTHER INFORMATION: GLUT4EX11 930 US-09-304-232-511
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 09-304-232-697
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LENGTH: 29
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US-08-454-557C-6/c

RESULT 90

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Patent No. 5948634
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: de la Monte, Suzanne
APPLICANT: de la Monte, Suzanne
APPLICANT: Wands, Jack R.
TITLE OF INVENTION: of Alzheimer's Disease
NUMBER OF SEQUENCES: 121
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox P.L.L.C.
STREET: 1100 New York Avenue, Suite 600
CITY: Washington
Patent No. 5830670
GENERAL INFORMATION:
APPLICANT: de la Monte, Suzanne
APPLICANT: de la Monte, Suzanne
APPLICANT: Wands, Jack R.
TITLE OF INVENTION: Neural Thread Protein Gene Expression and Detection
TITLE OF INVENTION: of Alzheimer's Disease
NUMBER OF SEQUENCES: 121
CORRESPONDENCE ADDRESS:
ADDRESSEE: Sterne, Kessler, Goldstein & Fox P.L.L.C.
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                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Sterne, Kessler, Goldstein & Fox P.L.L.C.
STREET: 1100 New York Avenue, Suite 600
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: D.C.
CUNDARY: U.S.A.
ZIP: 2005-3934
COMPUTER READABLE FORM:
MEDIUW TYPE: FIDOPY disk
COMPUTER: IBM PC Compatible
OPERATIOS SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/454,557C
FILING DATE: 30-MAY-1995
CLASSIFICATION: 514
ATTORNEY/AGERY INFORMATION:
NAME: LIDAWIG Steven R:
REGISTRATION NUMBER: 36,203
REPRENCE/DOCKET NUMBER: 36,203
REPRENCE/DOCKET NUMBER: 36,203
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPRAK: (202) 371-2600
INFORMATION FOR SEQ ID NO: 6:
SEQUENCE CHARACTERISTICS:
MANDE: LIDAMINE SEQUENCE CHARACTERISTICS:
MANDE: LIDAMINE SEQUENCE CHARACTERISTICS:
MANDE: LIDAMINE SEQ ID NO: 6:
MANDE: LIDAMINE SEQUENCE CHARACTERISTICS:
MANDE: LIDAMINE SEQUENCE C
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CURRENT APPLICATION DATA:
APPLICATION VMBER: US/08/340,426D
FILING DATE: 14-NOV-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 22.6; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pred. No. 2.1e+02;
0; Mismatches 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2179 TTCGCACCATTCTCCTGCCTCAGCCTCCC 2207
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COUNTRY: U.S.A.
ZIP. 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPALIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 86.2%;
Matches 25; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            nucleic acid
EDNESS: both
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS:
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CITY: Newport Beach
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STRANDEDNESS: both
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-579-445-10/c
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BASULT 8

BASULT 8

BACOL 450-673C-6/C

Sequence 6, Application US/08450673C

Patent No. 5948ATION.

APPLICANT: Wanda, Jack N.

TITLE OF INVENTION: Of Alzheimer's Disease

NUMBER OF SEQUENCES: 121

CORRESPONDERCE ADDRESS: 122

CONTRY: U.S.A.

STREET: 1100 New York Avenue, Suite 600

CITY: Washington

STREET: 100 New York Avenue, Suite 600

CITY: Washington

STREET: 100 New York Avenue, Suite 600

CITY: Washington

STREET: 100 New York Avenue, Suite 600

CITY: Washington

STREET: 100 New York Avenue, Suite 600

CITY: Washington

STREET: 100 New York Avenue, Suite 600

STREET: 100 Now Work Avenue, Suite 600

STREET: 100 Now Work Avenue, Suite 600

STREET: 100 Now Work Avenue, Suite 600

SCHWARE: Deather Norman Avenue, Marken 100

SERATION STEEM: PC-DOS/MS-DOS

SCHWARE: Deather Norman Avenue, Marken 100

SECUENCE CHARACTERISTICS:

LEEPENA: (202) 311-2540

INPORMATION POR SEG 10 NO: 6:

SEQUENCE CHARACTERISTICS:

LEMEGH: 30 Dase paire

TYPE: nucleic acid

TYPE: nucleic acid

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1.0%; Score 22.6; DB 1; Length 30; 86.2%; Pred. No. 2.1e+02; tive 0; Mismatches 4; Indels
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CLASSIFICATION: 435
ATTORNEY AGENT INFORMATION:
NAME: Ludwig, Steven R.
REGISTRATION NUMBER: 36,203
REFERENCE/DOCKET NUMBER: 0669.
TELECOMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
TELEPHONE: (202) 371-2540
TELEPHONE: (202) 371-2540
INFORMATION FOR SEQ ID NO: 6: SEQUENCE CHARACTERISTICS:
LENGTH: 30 base pairs
LENGTH: 30 base pairs
TYPE: mucleic acid
STRANDEDNESS: both
TOPOLOGY: both
US-08-340-426D-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 86.2
Matches 25, Conservative
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Matches 25; Conservative
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US-08-450-673C-6
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PETINGS 2.71111.6

PETINGS 2.7111.6

PETINGS C. PRUNKTION: Neural Thread Protein Gane Expression and TITLE OF INVENTION: Detection of Alzheimer's Disease NUMBER OF SEQUENCES: 12.

PETINGS PETINGS C. PETINGS 2.12

PETINGS PETINGS C. PETINGS 2.12

PETINGS PETINGS C. PETINGS 2.12

PETINGS PETINGS 2.12

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29 GATGGGGTTTCACCRTGTTGGCCAGGTTG 1
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LENGTH: 29
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Sequence 183, Application US/09304232

Batent No. 6552185

GENERAL INFORMATION:
APPLICANT: Fan, Jian Bing
APPLICANT: Chakravarti, Aravinda
APPLICANT: Glas Western Reserve University School of Medicine
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: 1990-05-03

CURRENT PAPLICATION NUMBER: US/09/304,232

CURRENT FILING DATE: 1999-05-03

EARLIER APPLICATION NUMBER: US 60/084,641

EARLIER APPLICATION NUMBER: US 60/084,641

EARLIER PETING DATE: 1999-05-03

SOUTHARE OF SEQ ID NOS: 2009

SOUTHARE PRACES FRASE FRASE FOR WINDOWS VERSION 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.9%; Score 22.4; DB 1; Length 25; Best Local Similarity 95.8%; Pred. No. 2.2e+02; Matches 23; Conservative 0; Mismatches 1; Indels
                                                                                               SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/579,445
FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/152,484
FILING DATE:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                   NAME: Kirkpatrick, Anita M.
REGISTRATION NUMBER: 32,617
REFERENCET DOCKET NUMBER: STRATAG.009A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 235-8550
TELEPAX: (619) 235-0176
INFORMATION FOR SEQ ID NO: 10: SEQUENCE CHARACTERISTICS:
LENGTH: 25 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2260 TTTTAGTAGAGACAGGGTTTCACC 2283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       24 TTTTAGTAGAGACAGATTCACC 1
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA ORGANISM: Artificial Sequence
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US-09-304-232-183
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HYPOTHETICAL: NO
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IMMEDIATE SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-304-232-183/c
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LENGTH: 29
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US-09-304-232-210/c

Sequence 210, Application US/09304232

Patent No. 65251805

GENERAL INFORMATION:

APPLICANT: Fan, Jian Bing

APPLICANT: Chakrawarti, Aravinda

APPLICANT: Affymetrix, Inc.

TITLE OF INVENTION: Polymorphisms Associated With

TITLE OF INVENTION: Hypertension

TITLE OF INVENTION: Hypertension

FILE REFERENCE: 018547-034210US

CURRENT APPLICANTON NUMBER: US/09/304,232

CURRENT APPLICATION NUMBER: US/09/404,232

CURRENT PILING DATE: 1999-05-03

EARLIER PILING DATE: 1998-05-07

NUMBER OF SEQ ID NOS: 909

SOUTHARD FALSE PASSEQ FOR Windows Version 3.0

LENGTH: 29
                                                                         GENERAL INFORMATION:
APPLICANT: Fan, Jian Bing
APPLICANT: Fan, Jian Bing
APPLICANT: Charkavarti, Aravinda
APPLICANT: Charkavarti, Aravinda
APPLICANT: Hubaka, Marc Kenneth
APPLICANT: Affewertix, Inc.
ATILE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Hypertension
FILE REFERENCE: 018547-034100S
CURRENT FILING DATE: 1999-05-03
EARLIER APPLICATION NUMBER: US 60/084,641
EARLIER APPLICATION NUMBER: US 60/084,641
EARLIER PILING DATE: 1999-05-03
NUMBER OF SEQ ID NOS: 909
SOFTWARE: FRESERE FOR WINDOWS Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indela
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0.9%; Score 22.2; DB 1;
Best Local Similarity 82.8%; Pred. No. 2.2e+02;
Matches 24; Conservative 1; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match

0.9%; Score 22.2; DB 1;
Best Local Similarity 82.8%; Pred. No. 2.2e+02;
Matches 24; Conservative 1; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2222 ACAGTCATCTGCCACCACACCTGGCTAAT 2250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2335 GCCTCCCAAAGTGCTGGGATTACAGGCAT 2363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  29 ACAGGCATCTGCCAYCATGCCCGGCTAAT 1
US-09-304-232-209/c
; Sequence 209, Application US/09304232
; Patent No. 6525185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: APOC4 2345
US-09-304-232-209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: APOC4 2366
US-09-304_232-210
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29 GCCTCCCGAGTAGCWGGGATTACAGGCAT 1

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TYPE: DNA
ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.9%
Best Local Similarity 82.8%
Matches 24; Conservative
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LENGTH: 29
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Sequence 696, Application US/09304232

Patent No. 6525185

GENERAL INFORMATION:

APPLICANT: Fan, Jian Bing

APPLICANT: Case Western Keserve University School of Medicine

APPLICANT: Affamerix, Inc.

TITLE OF INVENTION: Polymorphisms Associated With

TITLE OF INVENTION: Polymorphisms

APPLICANT: Associated With

APPLICANT: Associ
                                                                                                           GENERAL INFORMATION:

APPLICANT: Charavinia
APPLICANT: Charavinia
APPLICANT: Charavinia
APPLICANT: Case Western Reserve University School of Medicine
APPLICANT: Gase Western Reserve University School of Medicine
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: POlymorphisms Associated With
TITLE OF INVENTION: Hypertension
TITLE OF INVENTION: 1999-05-03
CURRENT APPLICATION NUMBER: US 60/084,641
EARLIER PILING DATE: 1999-05-07
NUMBER OF SEQ ID NOS: 909
SOFTWARE: FastSEQ for Windows Version 3:0
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                                      Sequence 686, Application US/09304232
Patent No. 6525185
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OTHER INFORMATION: PGISEX10 1505
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US-09-304-232-696
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Artificial Sequence
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Matches 24; Conservative
US-09-304-232-686/c
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LENGTH: 29
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                                        GENERAL INFORMATION:

APPLICANT: Fan, Jian Bing
APPLICANT: Chakravarti, Aravinda
APPLICANT: Chakravarti, Aravinda
APPLICANT: Halushka, Marc Kenneth
APPLICANT: Halushka, Marc Kenneth
APPLICANT: Affymetrix, Inc.
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Hypertension
FILE REFERENCE: 018547-03421005
CURRENT APPLICATION NUMBER: US/09/304,232
CURRENT APPLICATION NUMBER: US 60/084,641
EARLIER FILING DATE: 1998-05-07
NUMBER OF SEQ ID NOS: 909
SOFTWARE: FastSEQ for Windows Version 3.0
IENGTH: 29
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GENERAL INFORMATION:
APPLICANT: Fan, Jian Bing
APPLICANT: Chakravarti, Aravinda
APPLICANT: Chakravarti, Aravinda
APPLICANT: Halushka, Marc Kenneth
APPLICANT: Gae Western Reserve University School of Medicine
TITLE OF INVENTION: Polymorphisms Associated With
TITLE OF INVENTION: Hypertension
FILE REFERENCE: 018547-034210US
CURRENT FILING DATE: 1999-05-03
EARLIER APPLICATION NUMBER: US/09/304,232
CURRENT FILING DATE: 1999-05-03
EARLIER FILING DATE: 1999-05-03
NUMBER OF SEQ ID NOS: 909
SOFTWARE: PRESERVE FOR Windows Version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.9%; Score 22.2; DB 1;
82.8%; Pred. No. 2.2e+02;
tive 1; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2193 CTGCCTCAGCCTCCCAATTAGCTTGGCCT 2221
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Sequence 700, Application US/09304232 Patent No. 6525185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 863, Application US/09304232
Partent No. 6525185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; FEATURE:
; OTHER INFORMATION: PGISEX10 3061
US-09-304-232-700
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US-09-304-232-863
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Gaps
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## Sequence 3, Application US/09385917

## Patent No. 6294546

## APPLICANT: Rosen, Glenn

## APPLICANT: Russer, John

## ITILE OF INVENTION: ANTI-PROLIFERATIVE AGENT

## TITLE OF INVENTION: ANTI-PROLIFERATIVE AGENT

## TITLE OF INVENTION: ANTI-PROLIFERATIVE AGENT

## TITLE OF INVENTION: ANTI-PROLIFERATIVE AGENT

## CURRENT APPLICATION NUMBER: US/09/385,917

## CURRENT FILING DATE: 1999-08-30

## WUMBER OF SEQ ID NOS: 8

## SEQ ID'NO: 8

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20001 Jefferson Davis Highway, Suite 207
                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 1.44 megabyte, 3.5" microdisk
COMPUTER: Twinhead* Slimnote-890TX
OPERATING SYSTEM: MS DOS version 6.2,
OPERATING SYSTEM: Windows version 3.11
SOFTWARE: Word for Windows version 2.0
SOFTWARE: Converted to ASCI
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/100,104
FILING DATE:
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                                                                                         Virginia
: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2338 TCCCAAAGTGCTGGGATTACAG 2359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFCATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/107,673
FILING DATE: 18-Aug-93
APPLICATION NUMBER: 08/392,019
FILING DATE: 21-Feb-95
APPLICATION NUMBER: 08/571,047
FILING DATE: 12-Dec-95
APPLICATION NUMBER: 08/575,191
FILING DATE: 20-Dec-95
APPLICATION NUMBER: 08/575,191
FILING DATE: 22-Apr-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Friedmam, Mark M.
REGISTRATION NUMBER: 33,883
REFERENCE/DOCKET NUMBER: 205/15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 rcccaaagrecreedarracae 22
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TELEPHONE: 972-3-562553
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 972-3-5625554
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; STRANDEDNESS: singl
; TOPOLOGY: linear
US-09-100-104-1
                                          CITY: Ar
STATE: V
COUNTRY:
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                                                                                         Sequence 1, Application US/08635820A

Sequence 1, Application US/08635820A

Patent No. 5817462

GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION:
TITLE OF INVENTION:
WUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Mark M. Friedman c/o Robert Sheinbein
STREET: 2940 Birchtree lane
CITY: Silver Spring
STATE: Maryland
CONNTRY: United States of America
ZIP: 20906
COMPUTER READABLE FORM:
MEDDIUM TYPE: 1.44 megabyte, 3.5" microdisk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
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100.0%; Pred. No. 2.5e+02;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: 1.44 megabyte, 3.5" microdisk COMPUTER: 1.44 megabyte, 3.5" microdisk COMPUTER: Twinhead* Slimnote-890TX OPERATING SYSTEM: MS DOS veraion 6.2, OPERATING SYSTEM: Mindows version 3.11 SOFTWARE: Word for Windows version 3.11 SOFTWARE: CONVERTED VOICE WINDOWS VERSION 3.11 SOFTWARE: CONVERTED VOICE WINDOWS VERSION 3.11 APPLICATION NUMBER: US/08/635,820A FILING DATE: 22-Apr-1996 CLASSIFICATION NUMBER: 08/107,673 APPLICATION NUMBER: 08/107,673 APPLICATION NUMBER: 08/32,019 FILING DATE: 21-Feb-95 APPLICATION NUMBER: 08/571,047 FILING DATE: 12-Dec-95 APPLICATION NUMBER: 08/571,047 FILING DATE: 21-Dec-95 APPLICATION NUMBER: 08/571,047 FILING DATE: 20-Dec-95 APPLICATION NUMBER: 08/571,047 FILING DATE: 20-Dec-95 APPLICATION NUMBER: 08/575,191 FILING DATE: 20-Dec-95 APPLICATION NUMBER: 08/575,191
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Patent No. 6066459
GENERAL INFORMATION:
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REGISTRATION NUMBER: 33,883
REFERENCE/DOCKET NUMBER: 205
TELECOMMUNICATION INFORMATION:
TELEPHONE: 972-3-562553
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.'
Matches 22, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TOPOLOGY: linear
US-08-635-820A-1
                                                                                    -08-635-820A-1
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US-09-526-193A-275
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US-09-540-699-19
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Facent No. 6337984
GENERAL INFORMATION:
APPLICANT: Rosen, Glenn D.
APPLICANT: Lennox, Edwin S.
TITLE OF INVENTION: USES OF DITERPENOID TRIEPOXIDES AS AN TITLE OF INVENTION: ANTI-PROLIFERATIVE AGENT
FILE REFERENCE: STANO96DIV
CURRENT APPLICATION NUMBER: US/09/884,898
CURRENT PAPLICATION NUMBER: 09/385,917
PRIOR APPLICATION NUMBER: 09/385,917
PRIOR FILING DATE: 1999-08-30
NUMBER OF SEQ ID NOS: 8
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 3
LENGTH: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.9%; Score 22; DB 1; Length 22; Best Local Similarity 100.0%; Pred. No. 2.5e+02; Matches 22; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.9%; Score 22; DB 1; Length 22; Best Local Similarity 100.0%; Pred. No. 2.5e+02; Matches 22; Conservative 0; Mismatches 0; Indels
                         0; Indels
                                                                                                                                                                                                                    Sequence 19, Application US/09540699
Patent No. 6383752
GENERAL INFORMATION:
APPLICANT: Agrawal, Sudhir
APPLICANT: Kandimalla, Ekambar R.
TITLE OF INVENTION:
FILE REFERENCE: 99,128-B
CURRENT FILING DATE: 2000-03-31
PRIOR APPLICATION NUMBER: US/09/540,699
CURRENT FILING DATE: 2000-03-31
PRIOR PILING DATE: 1999-03-31
PRIOR APPLICATION NUMBER: US 60/127,138
PRIOR PILING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 26
NUMBER OF SEQ ID NOS: 26
100.0%; Pred. No. 2.5e+02; tive 0; Mismatches 0;
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                                                                   639 GTCAATCAGCAGGAATCATCGG 660
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                                                                                             1 GTCAATCAGCAGGAATCATCGG 22
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  Best Local Similarity 100.0
Matches 22; Conservative
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; ORGANISM: H. sapiens
US-09-884-898-3
                                                                                                                                                                               RESULT 105
US-09-540-699-19/c
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LENGTH: 22
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                                      ; GENERAL INFORMATION:
; APPLICANT: Rosen, Glenn D.
; APPLICANT: Rosen, Glenn D.
; APPLICANT: Lennox, Edwin S.
; TITLE OF INVENTION: Uses of Diterpenoid Triepoxides as an
; TITLE OF INVENTION: Anti-Proliferative Agent
; TITLE OF INVENTION: Anti-Proliferative Agent
; FILE REFERENCE: STANOSGURE: US/09/335,794
; CURRENT FILING DATE: 2001-08-22
; PRIOR APPLICATION NUMBER: 09/385,917
; PRIOR FILING DATE: 1999-08-30
; NUMBER OF SEQ ID NOS: 16
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 3
; LEMOTH: 22
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US-09-526-193A-275/c

Sequence 275, Application US/09526193A

Patent No. 6617122

GENERAL INFORMATION:
APPLICANT: Brooke-Wilson, Angela R.
APPLICANT: Brooke-Wilson, Angela R.
APPLICANT: Pimstone, Simon N.
TITLE OF INVENTION: CHOLESTEROL LEVELS

TITLE OF INVENTION: CHOLESTEROL LEVELS

TITLE OF INVENTION: CHOLESTEROL LEVELS

CURRENT APPLICATION NUMBER: 60/124,702

PRIOR APPLICATION NUMBER: 60/124,702

PRIOR FILING DATE: 1999-03-15

PRIOR FILING DATE: 1999-06-17

PRIOR PILING DATE: 1999-06-17

PRIOR PILING DATE: 1999-06-17

PRIOR PILING DATE: 1999-06-17

PRIOR FILING DATE: 1999-06-17

PRIOR FILING DATE: 1999-06-17

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 275

LENGTH. 27
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
0.9%; Score 22; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 2.5e+02;
Matches 22; Conservative 0; Mismatches 0;
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Best Local Similarity 100.0%; Pred. No. 2.5e+02;
Matches 22; Conservative 0; Mismatches 0;
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Sequence 3, Application US/09935794 Patent No. 6599499
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Homo sapiens
US-09-935-794-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Homo sapiens
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RESULT 112
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; Batent No. 6525185
; GENERAL INFORMATION:
; APPLICANT: Fan, Jian Bing
; APPLICANT: Case Western Kenneth
; APPLICANT: Case Western Keserve University School of Medicine
; APPLICANT: Affymetrix, Inc.
; TITLE OF INVENTION: Polymorphisms Associated With
; TITLE OF INVENTION: Pypertension
; TITLE OF INVENTION: Hypertension
; FILE REFERENCE: 018547-034210US
; CURRENT APPLICATION NUMBER: US 60/084,641
; EARLIER PELLING DATE: 1999-05-03
; NUMBER OF SEQ ID NOS: 909
; SOFTWARE FRENCE F
Sequence 181, Application US/09304232

Patent No. 6525185

GENERAL INFORMATION:
APPLICANT: Fan, Jian Bing
APPLICANT: Chakravarti, Aravinda
APPLICANT: Halushka, Marc Kenneth
APPLICANT: Alguentika, Marc Kenneth
APPLICANT: Affmetrix, Inc.
TITLE OF INVENTION: POlymorphisms Associated With
TITLE OF INVENTION: Pyertension
FILE REFERENCE: 018547-034210US
CURRENT FILING DATE: 1999-05-03
EARLIER APPLICATION NUMBER: US/09/304,232
CURRENT FILING DATE: 1999-05-03
BARLIER APPLICATION NUMBER: US 60/084,641
EARLIER OF FILING DATE: 1999-05-07
NUMBER OF SEQ ID NOS: 909
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US-09-304-232-509
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: APOC3 1736
US-09-304-232-181
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LENGTH: 29
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US-09-155-758-8

RESULT 111

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GENERAL INFORMATION:
APPLICANT: DORNER, Friedrich
APPLICANT: BARRETT: No. 61657111
APPLICANT: BIBL, Johann
TITLE OF INVENTION: PROCESS FOR DISINTEGRATING NUCLEIC ACIDS
TITLE OF INVENTION: AND PREPARING BIOLOGICAL PRODUCTS OF GUARANTEED QUALITY
NUMBER OF SEQUENCES: 10
CORRESPONDENCE ADDRESS:
ADDRESSEE: FOLEY & LARDNER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ö
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                                                                                                                                                                                                                                                                                                                                                STATE: D.C.

COUNTRY: U.S.A.

ZIP: ZOO07-5109

COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
APPLICATION NUMBER: US/09/155,758
FILING DATE: 08-APR-1997
PRIOR APPLICATION NUMBER: MO PCT/AT97/00068
FILING DATE: 08-APR-1997
PRIOR APPLICATION NUMBER: AT A 629/96
FILING DATE: 09-APR-1997
PRIOR APPLICATION NUMBER: AT A 629/96
FILING DATE: 09-APR-1997
APPLICATION NUMBER: 33,715
REFERENCE/DOCKET NUMBER: 33,715
REFERENCE/DOCKET NUMBER: 33,715
REFERENCE/DOCKET NUMBER: 33,715
REFERENCE/OFFUTION INFORMATION:
TELLEPHONE: (202) 672-5399
INFORMATION FOR SEQ ID NO: 8:
SEGUENCE CHRRACTERISTICS:
TEMPATH. 28 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-849-701-11/c
; Sequence 11, Application US/08849701
; Patent No. 5922544
; GENERAL INFORMATION:
; APPLICANT: Mayal, Kiyoshi
; APPLICANT: Naitoh, Tsutomu
; APPLICANT: Yonekawa, Toshihiro
; TITLE OF INVENTION: Method of Cell Detection
; NUMBER OF SEQUENCES: 12
; CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 21.6; DB 1;
Pred. No. 2.4e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Knobbe, Martens, Olson & Bear
STREET: 620 Newport Center Drive 16th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2100 GAGACCGAGICTTGCTCTGTTACCCAGG 2127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 gagacagagrcrcgcrcrgrcgcccagg 28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "Primer"
US-09-155-758-8
Sequence 8, Application US/09155758
                                                                                                                                                                                                                                                                                            3000 K Street, N.W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 85.7%;
Matches 24; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                nucleic acid
EDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Newport Beach
                                                                                                                                                                                                                                                                                                                    Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic
STRANDEDNESS:
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Gaps
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                                                                                                                                                                                                              0.9%; Score 21.4; DB 1; Length 23; 95.7%; Pred. No. 2.6e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 30, Application US/09618166
Patent No. 6583112
GENERAL INPORMATION
GENERAL INPORMATION:
APPLICANT: Fu, Ying-Hui
Yu, Chang-En
Oshima, Junko
Mulligan, John T.
Schellenberg, Gerald D.
TITLE OF INVENTION: GENE AND GENE PRODUCTS RELATED TO
WERNER'S SYNDROME
                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Seed Intellectual Property Law Group
STREET: 701 Fifth Avenue, Suite 6300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Seattle
STATE: Washington
COUNTRY: USA
ZIP: 98104-7092
COMPUTER READABLE FORM:
WEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/618,166
FILING DATE: 17-211-2000
CLASSIFICATION: 4UNKNOWN>
ATTORNEY/AGENT INFORMATION:
NAME: MCMASTERS, David D.
REGISTRATION NUMBER: 240052.419C1
TELECOMMUNICATION:
TELECOMMUNICATION: (206) 622-4900
                                                                                                                                                                                                                                                            0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 30:
                                                                                                                                                                                                                                                                                                          2336 CCTCCCAAAGTGCTGGGATTACA 2358
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2336 CCTCCCAAAGTGCTGGGATTACA 2358
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (206) 682-6
INFORMATION FOR SEQ ID NO: 30:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NUMBER OF SECUENCES: 209
; TELEFAX: (206) 682-6031
; INFORMATION FOR ESQ ID NO: 3
; SEQUENCE CHARACTERISTICS:
; LENGTH: 23 base pairs
; TYPE: nucleic acid
; STRANNEDNESS: single
; TOPOLOGY: linear
US-08-781-891-30
                                                                                                                                                                                                                                       Best Local Similarity 95.7
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Best Local Similarity 95.7
Matches 22, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                   JS-09-618-166-30/c
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                                                                                                                                                                                                              Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Fu, Ying-Hui
APPLICANT: Yu, Chang-En
APPLICANT: Yu, Chang-En
APPLICANT: Wulligan, John T.
APPLICANT: Schellenberg, Gerald D.
APPLICANT: Schellenberg, Gerald D.
TITLE OF INVENTION: WERNER'S SYNDROME
TITLE OF INVENTION: WERNER'S SYNDROME
CORRESPONDENCE: SEED and BERRY LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: PC-LUCS/NS-LUCS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/781,891
FILING DATE: 27-DEC-1996
CLASSIFICATION: 800
ATTORNEY/AGENT INFORMATION:
NAME: No. 6090620tenburg Ph.D., Carol
REGISTRATION NUMBER: 39.317
REPERENCE/DOCKET NUMBER: 240052.419
TELECOMMUNICATION INFORMATION:
TELEFHONE: (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    3: SEED and BERRY LLP
6300 Columbia Center, 701 Fifth Avenue
                                                                                                                                                                             APILING DATE:
CLASSIFICATION: 435
PION APPLICATION DATA:
APPLICATION NUMBER: PCT/JP95/02734
FILING DATE: 27-DEC-1995
ATTORNEY/AGENT INFORMATION:
NAME: Altman, Daniel B
REGISTRATION NUMBER: 34,115
REGISTRATION NUMBER: 34,115
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2335 GCCCCCAAAGTGCTGGGATTAC 2357
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STREET: 6300 COlumbia Center, 701
CITY: Seattle
STATE: Wanhington
COUNTRY: USA
ZIP: 98104-7092
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                             OPERATING SYSTEM: DOS
SCHWARE: RESESQ VERSION 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/849,701
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    23 GCCTCCCAAAGTGCTCGGATTAC 1
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Patent No. 6090620
                     COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 23 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 95.7<sup>1</sup>
Matches 22; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 714-760-9502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TOPOLOGY: linear
US-08-849-701-11
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US-08-781-891-30/c
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STATE: IL COUNTRY: United States of America ZIP: 60606
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: nucleic acid
HYPOTHERICAL: NO
ANTI-SENSE: NO
                                           Query Match 0.9
Best Local Similarity 100.
Matches 21; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Chicago
US-08-753-147-28
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                                                  APPLICANT: DUFP, GORDON
APPLICANT: DUFP, GORDON
APPLICANT: KORNMAN, KENNETH
APPLICANT: KORNMAN, KENNETH
APPLICANT: VAN DIJK, SIMON
TITLE OF INVENTION: DIAGNOSTICS AND THERAPEUTICS FOR EARLY-ONSET MENOPAUSE
FILE REFERENCE: MSA-012.01
CURRENT APPLICATION NUMBER: US/09/632,657
CURRENT FILING DATE: 2000-08-04
NUMBER OF SEQ ID NOS: 30
SOFTWARE: PATCHLIN VET. 2.1
SEQ ID NO 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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Sequence 28, Application US/08753147

SETENT No. 5770372

GENERAL INFORMATION:

APPLICANT: Concannon, Patrick

TITLE OF INVENTION: Detection of Mutations in the Human ATM Gene
NUMBER OF SEQUENCES: 196

CORRESPONDENCE ADDRESS:
ADDRESSEE: Christensen O'Connor Johnson and Kindness
STREET: 1420 5th Avenue
CITY: Seattle
STATE: Washingtion
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: Description of Artificial Sequence: Synthetic; OTHER INFORMATION: oligonucleotide
US-09-632-657-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 98101-2347
COMPUTER READABLE FORM:
MEDIUM TYEE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/753,147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Sheiness, Diama K.
REGISTRATION NUMBER: 35,356
REFERENCE/DOCKET NUMBER: VMRC-1-9714
TELECOMMUNICATION INFORMATION:
TELEFAK: (206) 749-4387
TELEFAK: (206) 224 0779
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2350 GGGATTACAGCCATGAGCCACCG 2372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 GGGATTACAGGCGTGAGCCACCG 23
         Sequence 25, Application US/09632657
Patent No. 6730476
                                                                                                                                                                                                                                                                                                                                                                 ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JS-08-753-147-28/c
                                                                                                                                                                                                                                                                                                                                             TYPE: DNA
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                                                                                                                                                                                                                   18-09-073-567-22

Sequence 22, Application US/09073567

Patent No. 601379-073-

GENERAL INFORMATION:

APPLICANT: Jiandong Chen

APPLICANT: Sudir Agrawal

APPLICANT: Ruiwen Zhang

TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLECTIDES

NUMBER OF SEQUENCES: 49

CORRESPONDENCE ADDRESS:

ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff

STREET: 300 South Wacker Drive, 32nd Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       %30-073-567-44/c
; Sequence 44, Application US/09073567
; Patent No. 601378.
; GENERAL INFORMATION:
    APPLICANT: Jiandong Chen
; APPLICANT: Sudhir Agrawal
; APPLICANT: Ruiwen Zhang
; TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
  Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 21;
                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.9%; Score 21; DB 1; Len
Pred. No. 2.8e+02;
0.9%; Score 21; DB 1; Lo 100.0%; Pred. No. 2.8e+02; tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.9%; Score 21; DB Best Local Similarity 100.0%; Pred. No. 2.6 Matches 21; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FILING DATE:
CLASSIFICATION:
ATTORNEY, AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REPRENCE/POCKET NUMBER: 96,057-A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 913-0002
; INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERESTICS:
LENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
FILING DATE:
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US-09-687-67B-18/C

Sequence 18, Application US/09687637B

Sequence 18, Application US/09687637B

Sequence 18, Application US/09687637B

SEGUENT NOTE OF COLUMBER 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence: MDM2 mRNA OTHER INFORMATION: reverse primer-2; +415 to +435.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      .
0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-540-699-21/c

JGS-09-540-699-21/c

JGS-09-540-699-21/c

JGENERAL INFORMATION:

JAPPLICANT: Agrawal, Sudhir

APPLICANT: Agrawal, Sudhir

TITLE OF INVENTION: Pseudo-Cyclic Oligonucleobases

TITLE REFERENCE: 99,128-B

CURRENT FILING DATE: 2000-03-31

PRIOR APPLICATION NUMBER: US 60/127,138

PRIOR APPLICATION NUMBER: US 60/127,138

PRIOR APPLICATION NUMBER: US 60/127,138

PRIOR PILING DATE: 2000-01-05

NUMBER OF SEQ ID NOS: 26

SOFTWARE: PATENTIN VON S: 26

SOFTWARE: PATENTIN VON S: 26

LENGTH: 21

TUBUGHH: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.9%; Score 21; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 2.8e+02;
Matches 21; Conservative 0; Mismatches 0;
                             NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1545
INFORMATION FOR SEQ ID NO: 269:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
LENGTH: 21 base pairs
TYPE: Nucleic Acid
STRANDENNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             726 GTACAAGAGCTTCAGGAAGAG 746
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA ORGANISM: Artificial Sequence
             TIORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                        ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-269
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 269, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDMZ
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U.S.A.
ZIP: 08053
COMPUTER REALABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Law Offices of Jane Massey Licata
STRET: 66 East Main Street
STATE: Marlton
STATE: NJ
                                                                                                                                         COMPUTER: IEM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERPECT 6.0
CURRENT APPLICATION DATA:
FILING DATE: herewith
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 119
US-09-280-805-269
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US-09-073-567-44
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SOFTWARE: FastSEQ for Windows Version 4.0
                              SEQ ID NO 90
LENGTH: 22
                                                                                                                           FEATURE:
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APPLICANT: Lander, Eric S.
APPLICANT: Lander, Eric S.
APPLICANT: Lander, James S.
APPLICANT: Ireland, James S.
APPLICANT: Ireland, James S.
APPLICANT: Bolk, Stacey
I APPLICANT: Bolk, George O.
APPLICANT: Bolky, George O.
APPLICANT: Bolky, George O.
APPLICANT: Daley, George O.
APPLICANT: Daley, George O.
APPLICANT: Daley, George O.
APPLICANT: Daley, George O.
APPLICANT: PROGRET 2825.1027-001
CURRENT APPLICATION NUMBER: US/09/657,472
CURRENT FILING DATE: 2000-09-07
PRIOR FILING DATE: 2000-09-07
PRIOR FILING DATE: 2000-07-26
PRIOR FILING DATE: 2000-07-26
PRIOR FILING DATE: 2000-08-16
NUMBER OF SEQ ID NOS: 2551
SOFTWARRE: PASLEEQ for Windows Version 4.0
SEQ ID NO 503
                                                                                                                                                                                                                                                     Query Match 0.9%; Score 20.8; DB 1; Length 27; Best Local Similarity 91.7%; Pred. No. 2.7e+02; Matches 22; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.9%; Score 20.6; DB 1; Length 21; 95.2%; Pred. No. 3e+02; tive 1; Mismatches 0; Indels
                                                                                                                                                                                  ; OTHER INFORMATION: Artificially synthesized primer sequence US-09-687-637B-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
APPLICANT: Brunkow, Mary
APPLICANT: Proil, Sean
APPLICANT: Prosh-Infow, Mary
APPLICANT: Paeper, Bryan
APPLICANT: Stanhling-Hampton, Karen
TITLE OF INVENTION: METHODS FOR IDENTIFYING
TITLE OF INVENTION: GENOMIC DELETIONS
FILE REPREBRUE: 240083.515
CURRENT APPLICATION NUMBER: US/09/918,686
CURRENT FILING DATE: 2001-07-30
PRIOR FILING DATE: 1998-04-14
NUMBER OF SEQ ID NOS: 46
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                  2192 CCTGCCTCAGCCTCCCAATTAGCT 2215
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 503, Application US/09657472 Patent No. 6727063
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NUMBER OF SEQ ID NOS: 105
                                                                                                               TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 95.2
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          -09-918-686-90/c
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                                                                       SEQ ID NO 18
LENGTH: 27
                                                                                                                                                                  FEATURE:
                                                                                            LENGTH:
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                                                                                                                                          Query Match 0.9%; Score 20.4; DB 1; Length 22; Best Local Similarity 95.5%; Pred. No. 3e+02; Matches 21; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 20.4; DB 1; Length 22;
Pred. No. 3e+02;
0; Mismatches 1; Indels
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APPLICANT: Hayden, Michael R.

APPLICANT: Brooks-Wilson, Angela R.

APPLICANT: Brinstone, Simon N.

TITLE OF INVENTION: METHODS AND REAGENTS FOR MODULATING

TITLE OF INVENTION: CHOLESTEROL LEVELS

FILE REPERENCE: 50110/002005

CURRENT APPLICATION NUMBER: US/09/526,193A

CURRENT FILING DATE: 2000-03-15
                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 94, Application US/09918686

Patent No. 6475739

GENERAL INFORMATION:
APPLICANT: Brunkow, Mary
APPLICANT: Brealing Hampton, Karen
APPLICANT: Saeahling Hampton, Karen
TITLE OF INVENTION: GENOMIC DELETIONS
TITLE OF INVENTION: GENOMIC DELETIONS
FILE REFERENCE: 240083.515
CURRENT PAPLICATION NUMBER: US/09/918,686
CURRENT PILING DATE: 2001-07-30
NUMBER OF SEQ ID NOS: 105
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 94
LENGTH: 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION NUMBER: 60/124,702
PRIOR APPLICATION NUMBER: 60/138,048
PRIOR APPLICATION NUMBER: 60/138,048
PRIOR FILING DATE: 1999-06-08
PRIOR APPLICATION NUMBER: 60/139,600
PRIOR FILING DATE: 1999-06-17
PRIOR PELING DATE: 1999-06-17
NUMBER OF SEQ ID NOS: 287
                                                                                                                                                                                                                                              2187 ATTCTCCTGCCTCAGCCTCCCA 2208
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; Sequence 274, Application US/09526193A
; Patent No. 6617122
                                                                                                                                                                                                                                                                                         22 ATTCTCTTGCCTCAGCCTCCCA 1
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TYPE: DNA ORGANISM: Artificial Sequence
                                                        ; OTHER INFORMATION: PCR primer US-09-918-686-90
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: PCR primer US-09-918-686-94
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al Similarity 95.5%;
21; Conservative
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Best Local Similarity
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US-09-073-567-4
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                                                                                                                                                                                                                                                                                                                                                         Sequence 2, Application US/09073567
Patent No. 6013786
GENERAL INFORMATION:
APPLICANT: Jandong Chen
APPLICANT: Sudhir Agrawal
APPLICANT: Ruiwen Zhang
TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
                                                                                                                                         0.9%; Score 20.4; DB 1; Length 22; 95.5%; Pred. No. 3e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.38+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Pred. ....
SOFTWARE: FastSEQ for Windows Version 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: United States of America
ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: FC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
FILING DATE:
CLASSIFICATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REGISTRATION NUMBER: 37,147
REGISTRATION INFORMATION:
TELEPHONE: (312) 913-0001
TELEPHONE: (312) 913-0002
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 Date pairs
                                                                                                                                                                                                                      2188 TTCTCCTGCCTCAGCCTCCCAA 2209
                                                                                                                                                                                                                                                             22 Trerecrecerrasecrecea 1
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MOLECULE TYPE: nucleic acid
                                                                                                                                Query Match
Best Local Similarity 95.5'
Matches 21; Conservative
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Best Local Similarity 100.
Matches 20, Conservative
                                                          TYPE: DNA
CORGANISM: Homo sapiens
US-09-526-193A-274
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STRANDEDNESS:
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US-09-073-567-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: IL
                                                                                                                                                                                                                                                                                                                                            US-09-073-567-2
                  SEQ ID NO 274
LENGTH: 22
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Gaps
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GENERAL INFORMATION:

APPLICANT: Jiandong Chen
APPLICANT: Sudhir Agrawal
APPLICANT: Sudhir Agrawal
APPLICANT: Ruiwen Zhang
TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREFT: 300 South Wacker Drive, 32nd Floor
CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT: Jiandong Chen
APPLICANT: Sudhir Agrawal
APPLICANT: Sudhir Agrawal
APPLICANT: Sudhir Agrawal
APPLICANT: Ruiwen Zhang
TITILE OF INVENTION: MDD2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: Chicago
CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
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3.3e+02;
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                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: IL.
COUNTRY: United States of America
ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OFFIGURE: Microsoft Word 97
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      98,057-A
                                                                                                                                                                                                                                                                      United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE.

CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFREENCE/DOCKEY NUMBER: 98,05
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 913-0001
TELEFAX: (312) 913-0001
TELEFAX: (312) 913-0002
SEQUENCE CHARACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  695 CCTTGAAGGTGGGAGTGATC 714
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Patent No. 6013786
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: nucleic acid
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 20 base pai
TYPE: nucleic acid
STRANDEDNESS: both
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US-09-073-567-3
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0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; Live 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Chicago
STATE: IL
COUNTRY: United States of America
                                                                                                                                           357 ACCTCACAGATTCCAGCTTC 376
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIPTCATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,05
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 913-0001
TELEPHONE: (312) 913-0001
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    369 CCAGCTTCGGAACAAGAGAC 388
                                                                                                                                                                                   1 ACCTCACAGATTCCAGCTTC 20
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                                    Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE
                                                                                                                                                                                                                                                                                                                    RESULT 130
US-09-073-567-8
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US-09-073-567-9
   Query Match
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APPLICANT: Jiandong Chen
APPLICANT: Jiandong Chen
APPLICANT: Gudit Agrawal
APPLICANT: Ruiwen Zhang
TITLE OF INVENTION: MDM.-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSE: 49
CORRESPONDENCE ADDRESS:
STREET: 300 South Wacker Drive, 32nd Floor
CITY: Chicago
STATE: 1L
COUNTRY: United States of America
ZIP: 60606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Ouery Match. 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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                      CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,057-A
TELECHONICATION INFORMATION:
TELEPHONE: (312) 913-0001
TELEFAX: (312) 913-0002
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1018 TGGATCAGGATTCAGTTTCA 1037
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Patent No. 6013786
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: nucleic acid
HYPOTHETICAL: NO
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; ANTI-SENSE: NO
US-09-073-567-7
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US-09-073-567-7
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                                                                                                                                                                                                     Sequence 8, Application US/09073567
Patent No. 6013786
GENERAL INFORMATION:
APPLICANT: Sudhir Agrawal
APPLICANT: Sudhir Agrawal
APPLICANT: Ruiwen Zhang
ITILE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: MCDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 9, Application US/09073567
Patent No. 6013786
GENERALY INFORMATION:
APPLICANT: Jiandong Chen
APPLICANT: Sudhir Agrawal
APPLICANT: Ruiwen Zhang
TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.8%; Score 20; DB 1; Length 20;
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Sequence 11, Application US/09073567

Patent No. 6013786

GENERAL INCRNATION:
APPLICANT: Sudhir Agrawal
APPLICANT: Sudwen Zhang
TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 4
ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
CITY: Chicago
STATE: IL
                                                                                                                                                                                                                                                                                                           Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,057-A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 913-0001
TELEPRX: (312) 913-0001
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: IL.

COUNTRY: United States of America
ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                          1203 TCCTTAGCTGACTATTGGAA 1222
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                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: nucleic acid
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear
MOLECULE TYPE: nucleic acid
                        TELEFAX: (312) 913-0002
INFORMATION FOR SEQ ID NO: 10
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                        ; ANTI-SENSE: NO
US-09-073-567-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ANTI-SENSE: NO US-09-073-567-11
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US-09-073-567-11
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APPLICANT: Jiandong Chen
APPLICANT: Sudmir Agrawal
APPLICANT: Minwen Zhang
STRET: 300 South Wacker Drive, 32nd Floor
CITY: Chicago
STRET: 1L
COUNTRY: United States of America
ZIP: 6006
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IMP PC COMPUTER: STORY:
MEDIUM TYPE: Floppy disk
COMPUTER: IMP PC COMPATIBLE
COMPUTER: WACKER PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION NUMBER: US/09/073,567
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ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
CITY: Chicago
                                                                                                     ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
FILING DATE:
CLASSIFICATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 98,057-A
TELEPHONE: (312) 913-0001
TELEPHONE: (312) 913-0002
TELEPHONE: (312) 913-0002
TELEPHONE: (312) 913-0002
TELEPHONE: CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: MUCIECI acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REPERENCE/DCKET NUMBER: 98,057-A
TELECOMMUNICATION INFORMATION:
                                                                                              United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 132
US-09-073-567-10
; Sequence 10, Application US/09073567
; Patent No. 6013786
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: nucleic acid
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ANTI-SENSE: NO
                                                                                              COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-073-567-9
                                                CITY:
STATE:
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STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 300 Sc
CITY: Chicago
MEDIUM TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-073-567-14
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Sequence 14, Application US/09073567

Sequence 14, Application US/09073567

Parent No. 6013786

GENERAL INFORMATION:
APPLICANT: Sudhir Agrawal
APPLICANT: Sudhir Agrawal
ITILE OF INVENTION:
NUMBER OF SEQUENCES: 49

CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
STREET: 300 South Wacker Drive, 32nd Floor
CITY: Chicago
STATE: 1L

COUNTRY: United States of America

ZIP: 60606
                                                                                                                                            APPLICANT: Jiandong Chen
APPLICANT: Sudhir Agrawal
APPLICANT: Sudhir Agrawal
APPLICANT: Rindon Abrang
TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
CITY: Chicago
STATE: IL
COUNTRY: United States of America
ZIP: 60606
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                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,057-A
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                   Sequence 13, Application US/09073567
Patent No. 6013786
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             669 ACATCTGTGAGTGAGACAG 688
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (312) 913-0002
INFORMATION FOR SEQ ID NO: 13
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: nucleic acid HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
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US-09-073-567-13
                                                             US-09-073-567-13
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COMPUTER: INE SCREEN PROCESS
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APPLICANT: Jiandong Chen
APPLICANT: Sudhir Agrawal
APPLICANT: Ruiwen Zhang
APPLICANT: Ruiwen Zhang
TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
                                                                                                                                       Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.38+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 1.

CORRESPONDENCE ADDRESS:
ADDRESSES: MCDORNeol Bochnen Hulbert & Berghoff STREET: 300 South Wacker Drive, 32nd Floor CITY: Chicago
STATE: 11
COUNTRY: United States of America ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC Compatible OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: IBM PC Compatible OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
TELEPHONE: (312) 913-0001
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OF SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LINFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Pred. ....
                                                                                                                                                                                                                                     680 TGAGAACAGGTGTCACCTTG 699
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US-09-073-567-17
; Sequence 17, Application US/09073567
; Betent No. 6013786
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 16, Application US/09073567
Patent No. 6013786
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                685 ACAGGTGTCACCTTGAAGGT 704
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                   ) MOLECULE TYPE: nucleic acid

) HYPOTHETICAL: NO

; ANTI-SENSE: NO

US-09-073-567-15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity 100.0
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: linear
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ANTI-SENSE: NO
US-09-073-567-16
                                                                                                                                                                                                                                                                                                                                                                               US-09-073-567-16
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APPLICANT: aidation Grand
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APPLICANT: aidation Graval
APPLICANT: withen Education
ITIME OF INVENTION: MON2.specific Control of Control
ITIME OF INVENTION: MON2.specific Graval
APPLICANT: mited States of America
CONTRY: United States of America
STRIE: 110.
CONTRY: United States of America
CONTRY: United States of America
CONTRY: United States of America
STRIE: 110.
CONTRY: MATCHING, MONARE: US/09/073,567
FILED OFFICE TO CONTRY: UNITED STRIES TO CONTROL OFFI
CONTRY: MATCHING, MONARE: US/09/073,567
FILEDOMNING, MONARE: US/09/073,567
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Gaps

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GENERAL INFORMATION:
APPLICANT: Jiandong Chen
APPLICANT: Jiandong Chen
APPLICANT: Jiandong Chen
APPLICANT: Jiandong Chen
APPLICANT: Sudhir Agrawal
APPLICANT: Ruiwen Zhang
TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 21, Application US/09073567
; Fatent No. 6013786
; GENERAL INFORMATION:
    APPLICANT: Jiandong Chen
    APPLICANT: Sudhir Agrawal
    APPLICANT: Ruiwen Zhang
    TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
    NUMBER OF SEQUENCES: 49
    CORRESPONDENCE ADDRESS:
    ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 20;
                                           Indels
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                   Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: United States of America ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98,057-A
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                                                                                                                                                                                                                                                  ; Sequence 20, Application US/09073567; Patent No. 6013786
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,05
TELECOMMINICATION INFORMATION:
TELEFAX: (312) 913-0001
TELEFAX: (312) 913-0002
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARATERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 TGAACATTCAGGTGATTGGT 20
                                                                                                                                  1 AAGGACCTTGTACAAGAGCT 20
                                                                                       717 AAGGACCTTGTACAAGAGCT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; ANTI-SENSE: NO
US-09-073-567-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL:
                                                                                                                                                                                                      RESULT 141
US-09-073-567-20
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STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Jiandong Chen
APPLICANT: Sudhir Agrawal
APPLICANT: Sudhir Agrawal
APPLICANT: Ruiwen Zhang
TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
CITY: Chicago
STATE: IL
COUNTRY: United States of America
ZIP: 60606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 20;
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,057-A
TELECOMMUNICATION INFORMATION:
TELEFAX: (312) 913-0001
TELEFAX: (312) 913-0002
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIPICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REPERENCE/DOCKET NUMBER: 37,147
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 913-0001
TELERAX: (312) 913-0001
INFORMATION FOR SEQ ID NO: 19: SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Flowers
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             5-09-073-567-19
Sequence 19, Application US/09073567
Patent No. 6013786
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       709 GTGATCAAAGGACCTTGTA 728
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GTGATCAAAGGACCTTGTA 20
                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
MOLECULE TYPE: nucleic acid
HYPOTHETICAL: NO
MATI-SENSE: NO
US-09-073-567-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: nucleic acid HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Jiandmag Chen
APPLICANT: Jiandmag Chen
APPLICANT: Sudhir Agrawal
APPLICANT: Ruiwen Zhang
TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS;
ADDRESSEE: MCDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
CITY: Chicago
STATE: IL
COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                         ö
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                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                   Score 20; DB 1; Le
Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                              Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIPICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REFERENCE/DOCKET NUMBER: 98,057-A
REFERENCE/DOCKET NUMBER: 98,057-A
TELECOMMUNICATION INFORMATION:
TELEFAX: (312) 913-0001
TELEFAX: (312) 913-0002
INFORMATION FOR SEQ ID NO: 24:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                      1027 ATTCAGTTTCAGATCAGTTT 1046
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1038 GATCAGTTTAGTGTAGAATT 1057
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 24, Application US/09073567
Patent No. 6013786
                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 ATTCAGTTTCAGATCAGTTT 20
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                                                                                                                                                           TOPOLOGY: linear
MOLECULE TYPE: nucleic acid
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MOLECULE TYPE: nucleic acid HYPOTHETICAL: NO
     TELEFAX: (312) 913-0002
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 base pairs
                                                                                                        TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 20 base pai
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                      ; ANTI-SENSE: NO US-09-073-567-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           HYPOTHETICAL: I ANTI-SENSE: NO US-09-073-567-24
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
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Patent No. 6013786

GENERAL INFORMATION:
ABPLICANT: Jiandong Chen
APPLICANT: Sudhir Agrawal
APPLICANT: Sudhir Agrawal
APPLICANT: Ruiwen Zhang
ITILE OP INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCL:
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
CITY: Chicago
STREET: United States of America
COUNTRY: United States of America
COUNTRY: Eloppy disk
COMPUTER READABLE FORM:
MEDIUM TYER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
FILTER ADDRESSEE: Microsoft Word 97
CURRENT APPLICATION DATA:
FILTER ADDRESSEE: Microsoft Word 97
FILTER ADDRESSEE: Microsoft WORDSER: US/09/073, 567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
300 South Wacker Drive, 32nd Floor
                                                                                  ZIP: G0606

COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,057-A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 913-0002
TELEPHONE: (312) 913-0002
TELEPHONE: CASE DID NO: 21:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REPRENCE/DOCKET NUMBER: 98,057-A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 913-0001
                                                                             United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1003 ATTCAGGTGATTGGTTGGAT 1022
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: nucleic acid
                             Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; ANTI-SENSE: NO
US-09-073-567-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-073-567-23
                                                                             COUNTRY:
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CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,057-A
TELECOMMUNICATION INFORMATION:
TELEFAX: (312) 913-0001
TELEFAX: (312) 913-0002
INFORMATION FOR SEQ ID NO: 29:
SEQUENCE CHARACTERISTICS:
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COUNTRY: United States of America
ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: 1BM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
PILING DATE:
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
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TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ANTI-SENSE: YES
US-09-073-567-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-073-567-29/c
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| Sequence 28, Application US/09073567
| Sequence 28, Application US/09073567
| Patent No. 6013786
| GENERAL INFORMATION:
| APPLICANT: Sudhir Agrawal
| APPLICANT: Sudhir Agrawal
| APPLICANT: Sudhir Agrawal
| TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
| TOTRESONDENCE ADDRESS:
| ADDRESSES: ADD
                                                      Sequence 27, Application US/09073567
Fatent No. 6013786
GENERAL INFORMATION:
APPLICANT: Jiandong Chen
APPLICANT: Jiandong Chen
TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
INWHER OF SEQUENCES: A
CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
CITY: Chicago
STATE: IL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,057-A
TELECHONIS: (312) 913-0001
TELEFAX: (312) 913-0002
INFORMATION FOR SEQ ID NO: 27:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 60606
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: MICROSCHE WORD 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: IL. COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              481 TTGGCCAGTATATTATGACT 500
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: nucleic acid HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      linear
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US-09-073-567-27
                                  US-09-073-567-27/c
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Best Local Similarity 100.0%; Pred. No. ...
Matches 20; Conservative 0; Mismatches
                             COMPUTER: IBM PC compatible
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Sequence 29, Application US/09073567
Patent No. 6013786
GENERAL INFORMATION:
APPLICANT: Standong Chen
APPLICANT: Standong Chen
APPLICANT: Standong Chen
APPLICANT: Standong Chen
TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESONDENCE ADDRESS:
ADDRESSE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor

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US-09-073-567-32/c
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                                                                                                                                                                                                                                                                                                                           Sequence 30, Application US/09073567
Patent No. 6013786
GENERAL INFORMATION:
APPLICANT: Jiandong Chen
APPLICANT: Ruiwen Zhang
TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLECTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: MCDOnnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
                                                                                                         Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STATE: IL COUNTRY: United States of America ZIP: 6666
COMPUTER READBLE FORM: MEDLUM TYPE: Floppy disk COMPUTER: ENOR POOR STATE OF SOFTWARE: MCTOSOFT WOOD 97
CURRENT APPLICATION DATA: APPLICATION DATA: APPLICATION NUMBER: US/09/073,567
FILING DATE: CLASSIFICATION: ATTORNEY/AGENT INFORMATION: ATTORNEY/AGENT INFORMATION: REFERENCE/DOCKET NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,057-A TELEFONNE: (312) 913-0001
TELEFONNE: (312) 913-0002
TELEFONNE: (312) 913-0002
TELEFONNE: (312) 913-0002
TELEFAX: (312) 913-0002
SERQUENCE CHARACTERIFICS: LENGTH: 20 DASE DAILS
LENGTH: 20 DASE DAILS
                                                                                                                                                                                           1018 TGGATCAGGATTCAGTTTCA 1037
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         357 ACCTCACAGATTCCAGCTTC 376
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Patent No. 6013706
GENERAL INFORMATION:
APPLICANT: Jiandong Chen
                                                                                                                                                                                                                  20 TGGATCAGGATTCAGTTTCA 1
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nucleic acid
NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.8
Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 300 Sc
CITY: Chicago
                                                YES
        MOLECULE TYPE:
                         HYPOTHETICAL:
ANTI-SENSE: Y
US-09-073-567-29
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APPLICANT: Buthir Agraed

APPLICANT: Ruisman Zhang

TITLE OF INVENTION: MOUSE-SPECIFIC ANTISENSE CLICONUCLEOTIDES

NUMBERS OF SEGURES: 49

CORRESONERS DESCRIPCES: 49

CORRESONERS DESCRIPCES: 49

CORRESONERS DESCRIPCES: 49

COMPATES READABLE FOR WELL

STATE: This doct a martica

COMPATES READABLE FOR COMPATION

SURTH TYPE: PORPY disk

WORDING TYPE: TO COMPATION

APPLICATION NUMBER: 19, 70

COMPATES TO COMPATES

COMPATES TO COMPATE

MARSHITCATION NUMBER: 19, 20, 74

FELENANCONIC NUMBER: 19, 21, 20, 20

TELENANCONIC NUMBER: 19, 21, 20

TELENANCONIC NUMBER: 10, 20, 20

TELENANCONIC NUMBER: 10, 20, 20

TELENANCONIC NUMBER: 19, 21, 20

TELENANCONIC NUMBER: 10, 20, 20

TELENANCONIC NUMBER: 10, 20

TELENANCONIC NUMBER: 10

TELENANCONIC NUMB
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Wed Jan 26 12:47:30 2005

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0; Mismatches
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                                              1203 TCCTTAGCTGACTATTGGAA 1222
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  20; Conservative
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STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
Matches 20; Conserva
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US-09-073-567-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
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  Matches
                                                                                                                                                                  RESULT 152
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Patent No. 6013786

GENERAL INFORMATION:

APPLICANT: Sudhir Agrawal

APPLICANT: Ruiwen Zhang

TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES

NUMBER OF SEQUENCES: 9

CORRESPONDENCE ADDRESS:

ADDRESSE: McDonnell Boehnen Hulbert & Berghoff

STREET: 300 South Wacker Drive, 32nd Floor

CITY: Chicago

STATE: IL

COUNTRY: United States of America

ZIP: 60606

COMPUTER: BADABLE FORM:

MEDIUM TYPE: Ploppy disk

COMPUTER: BM PC Compatible

COMPUTER: NICOSOFT WORD 97

CURRENT APPLICATION DATA:

ATTORNEY/AGENT INFORMATION:

ATTORNEY/AGENT INFORMATION:

MAME: Greenfield, Michael S.

REGISTRATION NUMBER: 37,147

TELECOMMUTICATION INFORMATION:

TELECOMMUTEN: 37,147

TELECOMMUTEN: 37,147

TELECOMMUTEN: 37,147

TELECOMMUTEN: 37,12) 91,20101
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              NAME: Greenfeld, Michael S.
REGISTRATION NUMBER: 37,147
REPERENG/DOCKET NUMBER: 37,147
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 913-0001
TELEFAX: (312) 913-0001
TELEFAX: (312) 913-0002
INFORMATION FOR SEQ ID NO: 32:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: linear
MOLECULE TYPE: nucleic acid
HYPOTHETICAL: NO
ANTI-SENSE: YES
US-09-073-567-32
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          780 TCTACCTCATCTAGAAGGAG 799
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TELEPHONE: (312) 913-0002
INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   nucleic acid
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-073-567-33/c
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Gaps
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Sequence 34, Application US/09073567

Sequence 34, Application US/09073567

Parent No. 6013786

GENERAL INFORMATION:
APPLICANT: Sudhir Agrawal
APPLICANT: Sudhir Agrawal
TITLE OF INVENTON: MMM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: MODONNEAL BOehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
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| Sequence 35, Application US/09073567
| Pattent No. 6013708.
| GENERAL INFORMATION:
| APPLICANT: Jiandong Chen
| APPLICANT: Sudir Agrawal
| APPLICANT: Ruiwen Zhang
| TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
| NUMBER OF SEQUENCES: 49
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
| STREET: 300 South Wacker Drive, 32nd Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 20;
Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 20; DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,057-A
TELECOMUNICATION INFORMATION:
TELEPHONE: (312) 913-0001
TELEFAX: (312) 913-0002
INFORMATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: United States of America ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
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US-09-073-567-37/c

| Sequence 37, Application US/09073567
| Patent No. 6013786
| Patent No. 6013786
| GENERAL INFORMATION:
| APPLICANT: Jiandong Chen
| APPLICANT: Sudhir Agrawal
| APPLICANT: Ruiwen Zhang
| TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
| NUMBER OF SEQUENCES: 49
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
| STREET: 300 South Wacker Drive, 32nd Floor
| CITY: Chicago
| STRIEE: Chicago
| STRIEE: Chicago
| STRIEE: Chicago
                                                                                                                                                                                                                                                     Length 20;
                                                                                                                                                                                                                                                                                                 0; Indels
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                                                                                                                                                                                                                                                     Query Match 0.8%; Score 20; DB 1; Le Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.8%; Score 20; DB 1; Le Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,057-A
TELEPHONE: (312) 913-0001
TELEPHONE: (312) 913-0002
INFORMATION FOR SEQ ID NO: 37:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
RPLICATION NUMBER: US/09/073,567
FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: IL
COUNTRY: United States of America
ZIP: 60606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  680 TGAGAACAGGTGTCACCTTG 699
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                                                                                                                                                                                                                                                                                                                                                                       20 GTGAGTGAGAACAGGTGTCA 1
                                                                                           , vectody: linear

, MOLECULE TYPE: nucleic acid

; HYPOTHETICAL: NO

; ANTI-SENSE: YES

US-09-073-567-36
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: nucleic acid HYPOTHETICAL: NO
  INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   HYPOTHETICAL: NC;
ANTI-SENSE: YES
US-09-073-567-37
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; Sequence 36, Application US/09073567
; Patent No. 6013786
; GENERAL INFORMATION:
; APPLICANT: Jiandong Chen
; APPLICANT: Sudhir Agrawal
; APPLICANT: Ruiwen Zhang
; TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
; NUMBER OF SEQUENCES: 49
; CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.38+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff STREET: 300 South Wacker Drive, 32nd Floor CITY: Chicago STATE: 11.

CONNTRY: United States of America ZIP: 60606

COMPUTER: EDADABLE FORM:
MEDIUM TYPE: Floppy disk COMPUTER: IBM PC COMPATIBLE FORM: PC-DOS/MS-DOS SOFTWARE: Microsoft Word 97

CURRALTON NUMBER: US/09/073,567

FILLING DATE:
                                                        ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: GREENIELD, NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,057-A
TELECOMMUNICATION INFORMATION:
TELEFAX: (312) 913-0001
TELEFAX: (312) 913-0002
INPORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
STARMDENNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 99,057-A
TELEPHONE: (312) 913-0001
TELEPHONE: (312) 913-0001
                                              United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                669 ACATCTGTGAGTGAGACAG 688
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        / Devology: linear MOLECULE TYPE: nucleic acid HYPOTHETICAL: NO / ANTI-SENSE: YES US-09-073-567-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 ACATCTGTGAGTGAGACAG 1
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Chicago
                                            COUNTRY: U
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OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; ANTI-SENSE: YES
US-09-073-567-39
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COUNTRY: Un
ZIP: 60606
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 158
US-09-073-567-40/c
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i Sequence 39, Application US/09073567

j Patent No. 6013786

j RARERAL INFORMATION:

APPLICANT: Jiandong Chen

APPLICANT: Sudhir Agrawal

APPLICANT: Chance And Application Application Applicant Applicant Applicant Applicant Applicant Applicant Applicant Applicant Application App
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                                                                                                                                                   APPLICANT: Jiandong Chen
APPLICANT: Jiandong Chen
APPLICANT: Jiandong Chen
APPLICANT: Sudhir Agrawal
APPLICANT: Ruiwen Zhawal
APPLICANT: Ruiwen Zhang
TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 20;
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FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 37,147
REFERENCE/OCKET NUMBER: 37,147
REGISTRATION FOR SEQ ID NO: 38:
REQUENCE CHARACTERISTICS:
FRUGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              United States of America
                               US-09-073-567-38/c
; Sequence 38, Application US/09073567
; Patent No. 6013786
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         685 ACAGGTGTCACCTTGAAGGT 704
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    linear
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Sequence 40, Application US/09073567

Patent No. 6013786

GENERAL INFORMATION:
APPLICANT: Sudhir Agrawal
APPLICANT: Sudhir Agrawal
APPLICANT: Sudhir Agrawal
APPLICANT: MINNED SPECIFIC ANTISENSE OLIGONUCLECTIDES
UNMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
                                                                                                                                                                                                   98,057-A
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?: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    704 TGGGAGTGATCAAAAGGACC 723
                                                                  FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,05
TELECOMMUNICATION INFORMATION:
TELEPAX: (312) 913-0001
INFORMATION FOR SEQ ID NO: 39:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: NUCLEIC acid
STRANDEDNESS: both
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
ATTORREY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFRENCE/DOCKET NUMBER: 98,05
TELECOMMUICATION INFORMATION:
TELEPHONE: (312) 913-0002
INFORMATION FOR SEQ ID NO: 40:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 receaercarcaaaecacc 1
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ATTORNEY/AGENT INFORMATION:
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US-09-073-567-42
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Jiandong Chen
APPLICANT: Sudhir Agrawal
APPLICANT: Ruiwen Zhawal
APPLICANT: Ruiwen Zhang
TITLE OF INVENTION: MDMZ-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES:
ADDERSSEE:
ADDERSEE:
ADDERSSEE:
ADDERSEE:
ADDERSSEE:
ADDERSEE:
ADDERSSEE:
ADDERSSEE
                                                                                                                                                                       Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.38+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff STREET: 300 South Wacker Drive, 32nd Floor CITY: Chicago STATE: 1L COUNTRY: United States of America ZIP: 60606 COMPUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS SOFTWARE: Microsoft Word 97 CURRENT APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION UNMBER: US/09/073,567 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TILLING DAILS:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,057
TELECOMMUNICATION INFORMATION:
TELERAX: (312) 913-0001
TELERAX: (312) 913-0002
INFORMATION FOR SEQ ID NO: 41:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 41, Application US/09073567
Patent No. 6013786
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 160
US-09-073-567-42/c
Sequence 42, Application US/09073567
Patent No. 6013786
                                                                                                                                                                                                                                                                                                                                                       709 GTGATCAAAGGACCTTGTA 728
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                                                                                                                                                                                                                                                                                                                                                                                                                                         20 GIGATCAAAGGACCTIGIA 1
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APPLICANT: Jiandong Chen
APPLICANT: Sudhir Agrawal
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 100. Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
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US-09-073-567-41
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 159
US-09-073-567-41/C
                                        ; ANTI-SENSE:
US-09-073-567-40
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US-09-073-567-43/c
; Sequence 43, Application US/09073567
; Sequence 43, Application US/09073567
; Patent No. 6013786
; GENERAL INFORMATION:
; APPLICANT: Usuden Chen
; APPLICANT: Ruiwen Zhang
; TTLE OF INVENTION: MOM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
; NUMBER OF SEQUENCES: 49
; CORRESPONDENCE ADDRESS: ADDRESS: ADDRESS: ADDRESSE: ADDRESS: 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ;
APPLICANT: Ruiwen Zhang
TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,057-A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 913-0001
TELEPHONE: (312) 913-0001
TELEPHONE: (312) 913-0001
SEQUENCE CHARACTERISTICS:
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ZIP: 6060

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
                                                                                                                                                                                                                                                                                   SIATE: IL COUNTRY: United States of America
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COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity
Matches 20; Conserv
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CLASSIFICATION:
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1027 ATTCAGTTTCAGATCAGTTT 1046
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-073-567-45/c
; Sequence 45, Application US/09073567
; Patent No. 6013786
; GENERAL INFORMATION:
; APPLICANT: Standing Chen
APPLICANT: Standing Chen
; TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
; CORRESPONDENCE ADDRESS:
; ADDRESSE: MCDORNell Boehnen Hulbert & Berghoff
; STREET: 300 South Wacker Drive, 32nd Floor
; CITY: Chicago
; STATE: IL
; COUNTRY: United States of America
ZIP: 60606
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NAME: Greenfield, MICHAEL

REGISTRATION NUMBER: 37,147

REGISTRATION NUMBER: 98,057-A

REFERENCE/DOCKET NUMBER: 98,057-A

TELECOMMUNICATION INFORMATION:
TELEPHAK: (312) 913-0002

INFORMATION FOR SEQ ID NO: 43:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: linear
MOLECULE TYPE: nucleic acid
HYPOTHETICAL: NO

""" GENURE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REPERENCE/DOCKET NUMBER: 98,057-A
TELECOMMUNICATION INFORMATION:
TELEFHONE: (312) 913-0001
TELEFAX: (312) 913-0002
INFORMATION FOR SEQ ID NO: 45:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1003 ATTCAGGTGATTGGTTGGAT 1022
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TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                        ; ANTI-SENSE: YES
US-09-073-567-43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILING DATE:
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US-09-073-567-45
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US-09-073-567-47/C
; Sequence 47, Application US/09073567
; Sequence 47, Application US/09073567
; Patent No. 6013786
; GENERAL INFORMATION:
; APPLICANT: Ujandong Chen
; APPLICANT: Guiden Agrawal
; APPLICANT: Ruiwen Zhamal
; APPLICANT: Ruiwen Zhamal
; TITLE: OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
; NUMBER OF SEQUENCES: 49
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
; STREET: 100 South Wacker Drive, 32nd Floor
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                                                                                                                Sequence 46, Application US/09073567
Sequence 46, Application US/09073567
Patent No. 6013786
GENERAL INFORMATION:
APPLICANT: Jiandong Chen
APPLICANT: Sudhir Agrawal
TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFERENCE/DOCKET NUMBER: 98,057-A
TELECOMMUNICATION INFORMATION:
TELEPHONE: (312) 913-0001
TELEFAX: (312) 913-0002
INFORMATION FOR SEQ ID NO: 46:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: ILL
COUNTRY: United States of America
ZIP: 60606
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/073,567
FILING DATE:
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20 ATTCAGTTTCAGATCAGTTT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: nucleic acid
HYPOTHETICAL: NO
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                        STREET: 300 Sk
CITY: Chicago
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; ANTI-SENSE: YES
US-09-073-567-46
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Chicago
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                                  TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 3
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
           609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 GCACCGCGCGAGCT
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Yes
                                                                                                                                                                                                                                                                       Linear
                                                                                                                                                                                                                                                               ; TOPOLOGY: Line; ANTI-SENSE: Yes US-09-280-805-3
       TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY:
ANTI-SENSE:
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US-09-280-801-3/C

18-09-280-801-3/C

18-09-280-801-3/C

18-09-280-805-3/C

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERRECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Microsoft Word 97
CURRENT APPLICATION DATA:
ATTONE DATE:
CLASSIFICATION NUMBER: US/09/073,567
FILING DATE:
ATTONES/AGENT INFORMATION:
NAME: Greenfield, Michael S.
REGISTRATION NUMBER: 37,147
REFERENE/DOCKET NUMBER: 98,057-A
TELEPHONE: (312) 913-0001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Licate, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
STATE: IL
COUNTRY: United States of America
ZIP: 60606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  675 GTGAGTGAGAACAGGTGTCA 694
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: nucleic acid
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity 100.(
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   , ANTI-SENSE:
US-09-073-567-47
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0.8%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
     Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-280-805-4/c
; Sequence 4, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: NO. COUNTRY: U.S.A.

ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPREFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION: DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jan Masch 26, 1998
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
TELECOMMUNICATION INFORMATION:
Query Match
0.8%; Score 20; DB 1; L
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
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COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44
COMPUTER: IBM PC
OPPERTING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Herewith
CLASSIFICATION NUMBER: 09/048,810
FILING DATE: MATCH 69/048,810
FILING DATE: MATCH 26,1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 1SPH-0346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                147 ATTAGTGCGTACGAGCGCCC 166
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TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 7:
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Best Local Similarity 100.0
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: Linear
HANTI-SENSE: Yes
US-09-280-805-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U.S.A. ZIP: 08053
MEDIUM TYPE:
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Patent No. 6184212

GENERAL INFORMATION:

APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2

TITLE OF INVENTION: EXPRESSION

NUMBER OF SEQUENCES: 271

CORRESPONDENCE ADDRESS:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street

COTY: Mailton
                                                  Sequence 5, Application US/09280805
Fatent No. 6184212
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: MAILON
STATE: NJ
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. ...a 0; Indels
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                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICCATION:
RELICATION NUMBER: 09/048,810
FILING DATE: MATCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 12,577
REFERENCE/DOCKET NUMBER: 13,2,57
REFERENCE/DOCKET NUMBER: 15PH-0346
TELLECOMMUNICATION INFORMATION:
TELLECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 08053
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-5
                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U.S.A.
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             RESULT 167
US-09-280-805-5/ç
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100.0%; Pred. No. 3.3e+02;
Live 0; Mismatches 0; Indels
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| Sequence 7, Application US/09280805
| Sequence 7, Application US/09280805
| Patent No. 6184212
| GENERAL INFORMATION:
| APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia
| APPLICANT: Graham, Brett P. Monia
| TITLE OF INVENTION: EXPRESSION
| TITLE OF INVENTION: EXPRESSION
| NUMBER OF SEQUENCES: 271
| CORRESPONDENCE ADDRESS:
DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street CITY: Marlton
            COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WONDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIPICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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Sequence 9, Application US/09280805 Patent No. 6184212
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US-09-280-805-9
                                           GENERAL INFORMATION:
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ZIP: 08053
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                                                                                                                                                                   0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MATCH 26, 1998
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match

0.8%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE, DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                      181 GAGAGTGGAATGATCCCCGA 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               273 CTCCAAGCGCGAAAACCCCG 292
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Patent No. 6184212
                                                                                                                                                                                                                                                                              20 GAGAGTGGAATGATCCCCGA 1
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                 i LENGTH: 20 base pairs
i TYPE: Nucleic Acid
i STRANDEDNESS: Single
i TOPOLGY: Linear
i ANTI-SENSE: Yes
US-09-280-805-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
                                                                                                                                                                                         Best Local Similarity 100.0
Matches 20; Conservative
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Linear
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US-09-280-805-8/c
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US-09-280-805-9/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        STATE: N. COUNTRY:
                                                                                                                                                                     Query Match
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; Sequence 10, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: EXPRESSION
; TITLE OF INVENTION: EXPRESSION
; UNMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; CARESPONDENCE ADDRESS:
; STREET: 66 East Main Street
; CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata
STREET: Ge East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER REALDBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INPORMATION:
NAME: VACENT INPORMATION:
NAME: APPLICATION:
NAME: APPLICA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM: MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        295 TGGTGAGGAGCAGGCAAATG 314
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 regreadeaceaceaare 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC OPERATING SYSTEM:
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Gequence 13, Application US/09280805; Patent No. 6184212; GENERAL INFORMATION:
                                                                                                                                                                                       331 CTGTACCTACTGATGGTGCT 350
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SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
LENGTH: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20; Conservative
    STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
                         ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-280-805-13/c
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ANTI-SENSE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-280-805-12
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 11, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION:
EXPRESSION
TITLE OF INVENTION: EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRA.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
CAMPITER: IBM PC
CAMPITER: IBM PC
CAMPITER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE LAW Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                          CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: MACCh 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licate, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ISPH-0346
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OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
SURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
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CLASSIFICATION
CLASSIFICATION
PRIOR PEPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jame Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-03
TELECOMMUNICATION INFORMATION:
TELEPRANE: 609-810-1515
TELEPRANE: 609-810-155
INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  303 AGCAGGCAAATGTGCAATAC 322
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                                                                                                                                                                                                                                                                TELEPHONE: 609-810-1515
TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                           herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 20 base pairs
TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: Linear

ANTI-SENSE: Yes

US-09-280-805-10
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STATE: NJ
COUNTRY: U.S.A.
                                                             FILING DATE:
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US-09-280-805-11/c
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Length 20;
                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 12, Application US/09280805
Fatent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, M
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A.

ZIP: 08053

COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
OPREATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGRAT INFORMATION:
NAME: Licata, Jane Massey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 20; DB 1; L. Pred. No. 3.3e+02;
Ouery Match

0.8%; Score 20; DB 1; L
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
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100.0%; Pred. No. 3...
... 0; Mismatches
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REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION:
TELEPHONE: 609-810-1515
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0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02;
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Patent No. 6184212
GENERAL INFORMATION
GENERAL INFORMATION
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
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CITY: 1 10 STATE: NJ
COUNTRY: 0.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
COMPUTER: SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION INDMER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
APPLICATION BATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MATCH 26,1998
ATTORNEY/AGENT INFORMATION:
ANDMEY/AGENT INFORMATION:
ANDMEY/AGENT INFORMATION:
ANDMEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match

Query Match
Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
                                      CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 15PH-0346
TELECOMMUNICATION NIPORMATION:
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1545
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1381 TTGATGTTCCTGATTGTAAA 1400
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NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH:
TELECOMMUNICATION INFORMATION:
TELEFAX: 609-810-1515
INFORMATION FOR SEQ ID NO: 15:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                 herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: Linear;
ANTI-SENSE: Yes
US-09-280-805-14
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ANTI-SENSE: Yes
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APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
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APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: U$/09/280,805
FILING DATE: herewith
CLASSIFICATION NUMBER: U$/09/88,810
FILING DATE: March 26, 1998
ATPORNEY/AGENT INFORMATION:
PROJECTION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 132,257
RESTERENCE/DOCKET NUMBER: 132,257
REGISTRATION STORE NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Score 20; DB 1; L. Pred. No. 3.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.4
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1047 AGTGTAGAATTTGAAGTTGA 1066
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Patent No. 6184212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 AGTGTAGAATTTGAAGTTGA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; ANTI-SENSE: Yes
US-09-280-805-13
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US-09-280-805-14/c
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US-09-280-805-16/c
i Sequence 16, Application US/09280805
j Patent No. 6184212
j Patent No. 6184212
j Patent No. 6184212
j APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia
j TITLE OF INVENTION: EXPRESSION
j TITLE OF INVENTION: EXPRESSION
j CORRESPONDENCE ADDRESS:
j CORRESPONDENCE ADDRESS:
j CORRESPONDENCE ADDRESS:
j STREET: 66 East Main Street
                                      Query Match

0.8%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
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US-09-280-805-17/c
US-09-280-805-17/c
; Patent No. 6184212
; GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: APPLICANT: Arabam, Breet P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER KEADMALE FOUR:

MEDIUM TYPE: DISCETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: 1BM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELEPHONE: 609-810-1454
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARCTERISTICS:
LENGTH: ALL 20 base pairs
TYPE: Nucleic Acid
STRENDENS: Single
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                                                                                                                              1695 TTTACATGTGCAAAGAAGCT 1714
                                                                                                                                                       20 TTTACATGTGCAAAGAAGCT 1
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Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET: 66 East
CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-280-805-16
US-09-280-805-15
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Gaps
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US-09-280-805-18/c
; Sequence 18, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Exhance Set P. Monia
TITLE OF INVENTION: Expression
; TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
                                                                                                                                                          COUNTRY:

ZIP:
20053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A.

ZIP: 08053
COMPUTER READALE FORM:
COMPUTER READALE FORM:
COMPUTER: IBM PC
COMPUTER: IBM PC
COMPUTER: IBM PC
COMPUTER: IBM PC
COMPUTER: OF PORTION WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILLING DATE: herewith
CLASSIETCATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 20; DB 1; Le:
Pred. No. 3.3e+02;
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1785 TAGTTGACCTGTCTATAAGA 1804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REPERCE/DOCKET NUMBER: ISPR
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPAX: 609-810-1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 17: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs TYPE: Nucleic Acid STRANDEDNESS: Single
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Best Local Similarity 100.0%; Pi
Matches 20; Conservative 0;
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; ANTI-SENSE: Yes
US-09-280-805-17
                                                                                                                      Marlton
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Indels

; Pred. No. 3.3e+02; 0; Mismatches 0;

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Best Local Similarity 100.0%;
                                                                          20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U
ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-280-805-21/c
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INFORMATION F
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                                                                          Matches
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Sequence 19, Application US/09280805
Fatent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
STATE: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM COMPUTER: IBM COMPUTER: IBM COMPUTER: IBM COMPUTER: SYSTEM: WINDOWS 95
SOFTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
RROW APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELEPHONE: 609-810-1515
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGIGIRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-07
TELECOMMUNICATION INFORMATION:
TELEFPAX: 609-810-1515
TELEFPAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 18:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TYPE: ANTI-SENSE: Yes
US-09-280-805-18
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INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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Best Local Similarity
Matches 20; Conserva
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U
ZIP: 08053
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0.8%; Score 20; DB 1; Length 20;

Query Match

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                                                                                                                                                                                                       GENERAL INFORMATION:

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.

APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2

TITLE OF INVENTION: EXPRESSION

NUMBER OF SEQUENCES: 271

CORRESPONDENCE ADDRESS:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 21, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Creen J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 20; DB 1; L
100.0%; Pred. No. 3.3e+02;
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIPRICATION:
PRICR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTATION NUMBER: 32,527
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ISPH-0346
1934 TAGTGGAATAGTGAATACTT 1953
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                                                                                                                                                    Sequence 20, Application US/09280805
Patent No. 6184212
                                              20 TAGTGGAATAGTGAATACTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEPHONE: 609-810-1515
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FORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                / ANTI-SENSE: Yes
US-09-280-805-20
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Sequence 25, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION:
ATTILE OF INVENTION:
TITLE OF INVENTION:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
STREET: NJ
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 20; DB 1; L. Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                100.0%; Pred. No. 3.3 tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIPICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGBNT INFORMATION:
NAME: Licata, Jane Massey
RESIRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                ISPH-0346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2256 GTACTTTTAGTAGAGACAGG 2275
                              REFERENCE/DOCKET NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH.
TELECOMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPAX: 609-810-1515
TELEPAX: 609-810-1654
INFORMATION FOR SEQ ID NO: 22:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
TOPOLOGY: Linear
WATI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 GTACTTTTAGTAGAGACAGG 1
Licata, Jane Massey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.8%
Best Local Similarity 100.0
Matches 20; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Nucleic Acid
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US-09-280-805-25/c
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US-09-280-805-25
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Sequence 22, Application US/09280805

Patent No. 6184212

GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: APPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                      COMPUTER: IBM PC COMPUTER: 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC COMP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT: INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.8%; Score 20; DB 1; Le Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0;
          E: Law Offices of Jane Massey Licata
66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street CITY: Marlton STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2224 AGTCATCTGCCACCACCT 2243
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                                                                                                                                                                    ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TOPOLOGY: Linear; ANTI-SENSE: Yes
US-09-280-805-21
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ZIP: 08053
                                                                                                                                     U.S.A.
   ADDRESSEE: Law
STREET: 66 East
CITY: Marlton
                                                                                                    STATE: NJ
COUNTRY: U.
ZIP: 08053
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RESULT 184

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Gaps

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NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 34: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs TYPE: Nucleic Acid STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity 100. Matches 20; Conservative
                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TOPOLOGY: Linear; ANTI-SENSE: Yes US-09-280-805-34
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U.S.A.
                        U.S.A.
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100.0%; Pred. No. 3.38+02;
                                                                                                                      US-09-280-805-33/c.

Sequence 33, Application US/09280805

Patent No. 6184212.

GENERAL INFORMATION:

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: EXPRESSION

TITLE OF INVENTION: EXPRESSION

WUMBER OF SEQUENCES: 271

CORRESPONDENCE ADDRESS:

ADDRESSEE: Law Offices of Jane Massey Licata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 187
US-09-280-805-34/c
; Sequence 34, Application US/09280805
; Patent No. 6184212
; GRNERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Arabam, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
; TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES:
; CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
MEDIUM TYPE: DISKETTE, 9.5
                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     3: Law Offices of Jane Massey Licata
66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           100.0%; Pred. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: 1BM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WONDERFECT 6.0
CUBRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY GRENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
37 GCCCTGTGTGTCGGAAGA 56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     4 CCGCGCGAGCTTGGCTGCTT 23
                      20 GCCCTGTGTCTCGGAAAGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 33:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20 ccececeaecricecrer 1
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Matches 20; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: Linear
TI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                  U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Law
STREET: 66 East
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                       STATE: NJ
COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; ANTI-SENSE:
US-09-280-805-33
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Gaps
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i Sequence 35, Application US/09280805

j Patent No. 6194212

i GENERAL INFORMATION:

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.

APPLICANT: Graham, Brett P. Monia

ITILE OF INVENTION:

TITLE OF INVENTION: EXPRESSION

NUMBER OF SEQUENCES: 271

CORRESPONDENCE ADDRESSS:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street

CORRESPONDENCE MAILON

MAILON

MAILON

STREET: 67 East Main Street

CORRESPONDENCE MAILON

                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; iive 0; Mismatches 0; Indels
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                           SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
RETERENCE/DOCKET NUMBER: 1SPH-0346
TELECHOMUNICATION INFORMATION:
TELECHOMUNICATION 118FORMATION:
TELECHOMUNICATION 1454
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
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TELECOMMUNICATION INFORMATION:

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ZIP: 08053
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Patent No. 6184212

GENERAL INPORMATION:

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.

APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: EXPRESSION

INVENTION: EXPRESSION

NUMBER OF SEQUENCES: 271

CORRESPONDENCE ADDRESS:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street

COTTY: Marlton

STATE: Marlton

STATE: Marlton

STATE: Marlton

STATE: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 20; DB 1; Length 20, Pred. No. 3.38+02;
                                                                                                                                                                                                                                      Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FLING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1) Score 20; DB 1) 100.0%; Pred. No. 3.3 Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: Warch 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
                                                                                                                                                                                                                                                                                                                            20 GCTTCTGGGGCCTGTGTGGC 39
                                                                                                                                                                                                                                                                                                                                                                    20 GCTTCTGGGGCCTGTGTGGC 1
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INFORMATION FOR SEQ ID NO: 36:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
| TELEPHONE: 609-810-1515 |
| TELEFAX: 609-810-1515 |
| TELEFAX: 609-810-1544 |
| INFORMATION FOR SEQ ID NO: 35: SEQUENCE CHARACTERISTICS: LENGTH: 20 base parrs |
| TYPE: Nucleic Acid |
| TYPE: Nucleic Acid |
| STRANDEDNESS: Single |
| TOPOLOGY: Linear |
| ANTI-SENSE: Yes |
| US-09-280-805-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 20; Conservat
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                                                                      Sequence 37, Application US/09280805
; Sequence 37, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: EXPRESSION
; UNMER OF SEQUENCES: 27;
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
; CITY: Mailton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 20; DB 1; Length 20;
Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CTIT: ...

STATE: NJ
COUNTRY: U.S.A.

ZIP: 08053

ZIP: 08053

COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
COMPUTER: MORDERFECT 6.0

SOFTWARE: WORDERFECT 6.0

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805

FILING DATE: herewith
CLASSIFICATION: PRIOR APPLICATION ATA:
APPLICATION NUMBER: 09/048,810

FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTATION NUMBER: 32,257
REFERENCE/LOCKET NUMBER: 32,257
RECOMMUNICATION INFORMATION:
RECOMMU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 38, Application US/09280805
| Patent No. 6184212
| Patent No. 6184212
| Patent No. 6184212
| Patent No. 6184210
| Patent No. 6184210
| APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia
| TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
| TITLE OF INVENTION: EXPRESSION
| NUMBER OF SEQUENCES: 271
| CORRESPONDENCE ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: ADDRESS: STREET: Ge East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; bcc.
100.0%; Pred. No. ...
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 100.0
Matches 20; Conservative
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STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Linear
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US-09-280-805-37
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Marlton
RESULT 190
US-09-280-805-37/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 191
US-09-280-805-38/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: N. COUNTRY:
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Gaps

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US-09-280-805-40/c

US-09-280-805-40/c

Sequence 40, Application US/09280805

Patent No. 6184212

GENERAL INFORMATION:

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.

APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2

TITLE OF INVENTION: EXPRESSION

NUMBER OF SEQUENCES: 271

CORRESPONDENCE ADDRESS:

ADDRESSE: Law Offices of Jane Massey Licata

STREET: 65 East Main Street

STREET: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                          0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OPERATING SISTEM: MINDLOWS 30 SOFTWARE: WORDPERFECT 6.0 CURRENT APPLICATION DATA: APPLICATION UNMBER: US/09/280,805 FILING DATE: herewith CLASSIFICATION NUMBER: 09/048,810 FILING DATE: MARCH 26, 1998 APPLICATION NUMBER: 09/048,810 FILING DATE: MARCH 26, 1998 ATTORNEY/AGENT INFORMATION: NAME: Licata, Jane Massey REGISTRATION NUMBER: 32,257 REFERENCE/DOCKET NUMBER: 1SPH-0346 TELECOMMUNICATION INFORMATION: TELECOMMUNICATION INFORMATION: 609-810-1515 TELEFAX: 609-810-1454 INFORMATION FOR SEQ ID NO: 40: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs TYPE: Nucleic Acid STRANDENES: SINGle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    50 GGAAAGATGGAGCAAGAAGC 69
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              information for SEQ ID NO: 3 SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs TYPE: Nucleic Acid STRANDEDNESS: Single STRANDEDNESS: Single STRANDEDNESS: Single STRANDEDNESS: Yes US-09-280-805-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Linear
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US-09-280-805-40
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                                                                                                                                                                                                                                                                                                                                                 Query Match
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0.8%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 3.38+02;
Matches 20; Conservative 0; Mismatches 0; Indels
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Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Coren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: 27PRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
COMPUTER READABLE FORM:

MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION:
PRIOR APPLICATION:
PRIOR APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY, AGENT INFORMATION:
NAME: Licata, Jame Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 15PH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1545
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: NUMCL 21 Acid
STREET STREET SAIDS
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CITY: Mariton
STATE: NJ
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REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
FELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
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; ANTI-SENSE: Yes
US-09-280-805-38
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0.8%; Score 20; DB 1; Length 20;
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Patent No. 6184212
GENERAL INFORMATION
GENERAL INFORMATION
GENERAL INFORMATION
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: US.A.

ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYER: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
RICHARD APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MAZCH 26, 1998
ATTORNEY/AGENT INFORMATION:
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Best Local Similarity 100.0%; Pred. No. ...
Marches 20; Conservative 0; Mismatches
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                                       SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
                                                                                                             APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-034
TELECOMUNICATION INFORMATION:
TELECOMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          98 TGACCGAGATCCTGCTTT 117
      OPERATING SYSTEM: WINDOWS 95
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REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: IS
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Marlton
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US-09-280-805-42
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US-09-280-805-43/c
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                          Sequence 41, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: 271
CORRESPONDENCE: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Mailton
STATE: NJ
                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.4 COMPUTER: 1BM PC OPERATING SYSTEM: WINDOWS 95 SOFTWARE: WORDPERFECT 6.0 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/280,805 FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION: 10FORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
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; Sequence 42, Application US/09280805
Patent No. 6184212
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: Linear ANTI-SENSE: Yes
US-09-280-805-41/c
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GENERAL INFORMATION:
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                                                                                                                   Score 20; DB 1; Length 20;
Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                               Sequence 44, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
UNMER OF SEQUENCES: 271
CORRESSED IAW OFFICES OF Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                               Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: COMPUTER: IBM PC
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: herewith (CLASSIFICATION) FIREWITH CLASSIFICATION DATA: APPLICATION DATE: APPLICATION NUMBER: 09/048,810 FILING DATE: MARCH 26, 1998 ATTORNEY/AGENT INFORMATION: NAME: Licata, Jane Massey
                                                                                                                                                                                                 105 GATCCTGCTGCTTTCGCAGC 124
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; Sequence 45, Application US/09280805
; Patent No. 6184212
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REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISE
TELECOMMUNICATION: 609-810-1515
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TYPE: Nucleic Acid
STRANDEDNESS: Single
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Best Local Similarity
                                    TOPOLOGY: Linear
ANTI-SENSE: Yes
US-09-280-805-43
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U.S.A.
ZIP: 08053
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STATE: NJ
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Patent No. 6184212
GENERAL INFORMATION:
TIPEMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2 TITLE OF INVENTION: EXPRESSION NUMBER OF SEQUENCES: 271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STALE.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
CAMPUTER: IBM PC
THE STATEMENT OF THE STORAGE
THE STATEMENT OF T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIPICATION:
RRICH APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            120 GCAGCCAGGAGCACCGTCCC 139
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COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 609-810 INFORMATION FOR SEQ ID NO: 4 SEQUENCE CHARACTERISTICS:
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STRANDEDNESS: Single
TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      U.S.A.
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ZIP: 08053
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US-09-280-805-46/c
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                                                             Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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100.0%; Pred. No. 3.3e+02;
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                                                                                                                                                                                                                                                                                           Sequence 48, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION:
EXPRESSION
TITLE OF INVENTION:
EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IEM PC
OOMPUTER: IEM PC
OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 202..
US-09-280-805-49/c
; Sequence 49, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS: Law Offices of Jane Massey Licata STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          100.0%; Prec. ...
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAMME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TYPE: Nucleic Acid
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Matches 20; Conserva
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                                                                                                                                                                                                                                                      RESULT 201
US-09-280-805-48/c
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  ; ANTI-SENSE:
US-09-280-805-47
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Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE. DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
GURRENT APPLICATION DATA
APPLICATION NUMBER: US/09/280,805
FILING DATE:
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Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 Bast Main Street
CITY: Marlton
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                        PRIOR APPLICATION DATA:
APPLICATION NUMBER:
PILING DATE: MACCA 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey,
REGISTRATION NUMBER: 32,257
REPERENCE/DOCKET NUMBER: ISPH-0346
TELECHONE: 609-810-1515
TELEPHONE: 609-810-1515
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REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1454
APPLICATION NUMBER: US/09/280,805
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  150 AGTGCGTACGAGCGCCCAGT 169
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INPORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
INTPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                   herewith
                                                                                                                                                                                                                                                                                                                                                                            Linear
Yes
                   FILING DATE: h
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ANTI-SENSE:
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Length 20;
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Sequence 51, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
TITLE OF INVENTION: RYPRESSION
TITLE OF INVENTION: EXPRESSION
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
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MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MATCH 26, 1998
ATTORNEY/AGENT INFORMATION:
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CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
MPUTED: A.S.A.
MATHER OF THE STATE OF THE
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0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
PRIOR APPLICATION DATA:

APPLICATION NUMBER: 09/048,810
FILING DATE: Warch 26,1998
ATTORNEY, AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPAX: 609-810-1515
STRANDENNESS: SINGLE
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REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPACNE: 609-810-1454
INFORMATION FOR SEQ ID NO: 51:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
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; STRANDEDNESS: Single
; TOPOLOGY: Linear
US-09-280-805-51
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: Linear
ANTI-SENSE: Yes
US-09-280-805-50
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Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marten
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                           CITY: MAAA-STATE: NJ
COMPUTRY: U.S.A.
21P: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: BM PORTER: BM NOWS 95
SOFTWARE: WORDPERRECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION NUMBER: 09/048,810
FILING DATE: MATCA 26, 1998
ATTORNEY/AGENT INFORMATION:
BRICH APPLICATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 609-810-1516
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: NUCLOSY: Linear
TYPE: NUCLOSY: Linear
TYPE: NUCLOSY: Linear
       ANTISENSE MODULATION OF HUMAN MDM2 EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: TBM PC:
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
RAPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
   TITLE OF INVENTION: ANTISENSE MODULATION OF HUN
TITLE OF INVENTION: EXPRESSION
CORRESPONDENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Mailton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              174 TGGCCCGGAGAGTGGAATGA 193
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Best Local Similarity 100.0
Matches 20; Conservative
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ZIP: 08053
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| Sequence 52, Application US/09280805
| Patent No. 6194212|
| Patent No. 6194212|
| Patent No. 6194212|
| APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia
| TITLE OF INVENTION: ANTISRNSE MODULATION OF HUMAN MDM2 TITLE OF INVENTION: EXPRESSION NUMBER OF SEQUENCES: 271
| CORRESPONDENCE ADDRESS: 271
| CORRESPONDENCE ADDRESS: 371
| COUNTRY: U.S.A.
Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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; Patent No. 6184212
; GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Herewith
CLASSIFICATION: DATA:
APPLICATION DATA:
PRIOR APPLICATION
PRIOR APPLICATION
PRIOR APPLICATION
ANDER: MACCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REPERENCE/DOCKET NUMBER: 32,257
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                                                                                                                                                                                                     CCGCAGTA 227
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                                                                                                                                                                                                208 GCCGTCGTGCT1
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US-09-280-805-52
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US-09-280-805-54/C
; Sequence 54, Application US/09280805
; Retent No. 6184212
; GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
; TITLE OF INVENTION: EXPRESSION
; VUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jame Massey Licata
; STREET: 66 East Main Street
; STREET: Marlton
; STREET: Narlton
; STREET: Narlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02;
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ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                 ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
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OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIPETCATION:
PRIOR APPLICATION ONTA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26,1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INPORMATION:
TELEBORMONICATION INPORMATION:
TELEBORMONICATION INPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ISPH-0346
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    242 AGGAAACTGGGGAGTCTTGA 261
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 20 base pairs
TYPE: Nucleic Acid
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
                                                                                        STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
                                                                  Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-280-805-53
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293 GATGGTGAGGAGCAGGCAAA 312
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SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        609-810-1454
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US-09-280-805-56
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A.
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US-09-280-805-55/c
| US-09-280-805-55/c
| Sequence 55, Application US/09280805
| Patent No. 6184212
| GENERAL INFORMATION:
| APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2 | TITLE OF INVENTION: EXPRESSION | NUMBER OF SEQUENCES: 271
| CORRESPONDENCE ADDRESS: | ADDRESSE: Law Offices of Jane Massey Licata | STREET: 66 East Main Street | CITY: Mailton | CITY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.38+02; Matches 20; Conservative 0; Mismatches 0; Indels
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COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Licata, Jane Massey
REGIGSTRATION NUMBER: 32,257
REGIGSTERENCE/DOCKET WUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                    ISPH-0346
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      289 CCCGGATGGTGAGGAGCAGG 308
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION UNDRER: 32,257
REFERENCE/DOCKET UNDRER: ISPH-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPHOR: 609-810-1454
INFORMATION FOR SEQ ID NO: 54:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
US-09-280-805-54
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 CCCGGATGGTGAGGAGCAGG 1
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INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
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Best Local Similarity 100.
Matches 20; Conservative
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STATE: NJ
COUNTRY: U
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; ANTI-SENSE:
US-09-280-805-55
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US-09-280-805-56/c

is Squence 56, Application US/09280805

is Batent No. 6184212

GENERAL INFORMATION:

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.

APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2

TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2

TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2

TITLE OF INVENTION: ANTISENSE

CORRESPONDENCE ADDRESS:

ADDRESSE: Law Offices of Jane Massey Licata

STREET: 65 East Main Street

STREET: NJ

COTATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ACREAGE STATES APPLICATION US/09280805
; Sequence 57, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
    APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
    APPLICANT: Graham, Brett P. Monia
    TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
    TITLE OF INVENTION: EXPRESSION
    NUMBER OF SEQUENCES: 271
    CORRESPONDENCE ADDRESS:
    CORRESPECT ADDRESSE: Law Offices of Jane Massey Licata
    STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:

21P: 0804

21P: 0804

21P: 0804

COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: 1BM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: U$/09/280,805
FILING DATE: herewith
CLASSIPICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 32,257
REFERENCE/DOCKET NUMBER: TSPH-0346
TELLCOMMUNICATION INFORMATION:
TELLECOMMUNICATION INFORMATION:
TELLECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 20; DB 1; I
Pred. No. 3.3e+02;
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0.8%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 59, Application US/09280805; Sequence 59, Application US/09280805; Patent No. 6184212.
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2; TITLE OF INVENTION: EXPRESSION
INTMER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSER: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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0.8%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U.S.A.
ZIP: 08053
ZIP: 08053
COMPUTER READBLE FORM:
COMPUTER: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: WORDPERPECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION
APPLICATION:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENI INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
            REFERENCE/DOCKET NUMBER: ISPH-0346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                297 GTGAGGAGCAGGCAAATGTG 316
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                               TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1115
TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 58:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDENNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 GTGAGGAGCAGGCAAATGTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
TENGTH: 20 base pairs
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TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                            ; TOPOLOGY: Linear; ANTI-SENSE: Yes US-09-280-805-58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: Linear
NATI-SENSE: Yes
US-09-280-805-59
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US-09-280-805-59/c
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Patent No. 6184212

GENERAL INFORMATION:

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.

APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2

TITLE OF INVENTION: EXPRESSION

NUMBER OF SEQUENCES: 271

CORRESPONDENCE ADDRESS:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street

CITY: Mailton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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COUNTRY: U.S.A.

ZIP: 08053

COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: TBM PC.
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERPECT 6.0
CARRENT APPLICATION DATA:
CARRENT APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION SEQ 155
TELEFAX: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                296 GGTGAGGAGCAGCAAATGT 315
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TOPOLOGY: Linear; ANTI-SENSE: Yes
US-09-280-805-57
                                                               U.S.A.
         Marlton
                                                                                              08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-280-805-58/c
CITY: Mas
STATE: N.
COUNTRY:
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DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: NO COUNTRY: NO COUNTRY: NO COUNTRY: NO COUNTRY: NO COUNTRY: NEADDBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44
COMPUTER: IBM PC
COMPUTER: IBM PC
COMPUTER: IBM PC
COMPUTER: OS COTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTONNEY, AGBNT INPORMATION:
NAWE: LICATEA, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 300 AGGAGCAGGCAAATGTGCAA 319
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-280-805-62/c
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                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               3.3e+02;
hes 0; Indels
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Patent No. 6184212
GENERAL INFORMATION
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION OF HUMAN MDM2
TITLE OF HUMAN MDM2
TITLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTEX: 0.5.A.

ZIP: 08053
COMPUTEX: 0.5.A.

MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: DISKETTE, 0.5
CURRATING SYSTEM: WINDOWS 95
SOFTWARE: WORDBERFECT 6.0
CURRATING SYSTEM: WINDOWS 95
CURRATION NUMBER: U5/09/280,805
FILING DATE: herewith
CLASSIPICATION NUMBER: 09/048,810
FILING DATE: march 26, 1998
ATTORNEY, AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERRING DOCKET NUMBER: 15.PH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1155
TELEFRANCE COS-810-1155
TELEFRANCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: SINGle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match

0.8%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
                                                                                                                RESULT 213
US-05-280-805-60/c
'Sequence 60, Application US/09280805
'Patent No. 6184212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        299 GAGGAGCAGGCAAATGTGCA 318
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20 TGAGGAGCAGGCAAATGTGC 1
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COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Marlton
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION NUMBER: 09/048,810
FILING DATE: MATCA 186,1998
ATTORA APPLICATION NUMBER: 09/048,810
FILING DATE: MATCA 26,1998
ATTORA APPLICATION NUMBER: 1257
REFERENCE CAPACITY INFORMATION:
NAME: Liceta, Jane Massey
REGISTRATION NUMBER: 1257
REFERENCE COSTET NUMBER: 15PH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEFAK: 609-810-1515
TELEFAK: 609-810-155
TELEFAK: 20 base pairs
TENDORMATION FOR SEQ 1D NO: 61:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TENDORMATION FOR SEQ 1D NO: 61:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TENDORMATION FOR SEQ 1D NO: 61:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TENDORMATION FOR SEQ 1D NO: 61:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TENDORMATION FOR SEQ 1D NO: 61:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TENDORMATION FOR SEQ 1D NO: 61:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TENDORMATION FOR SEQ 1D NO: 61:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TENDORMATION FOR SEQ 1D NO: 61:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
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3.3e+02;
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US-09-280-805-65/c
                      US-09-280-805-64/C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 20; DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDMZ
TITLE OF INVENTION: EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       E: Law Offices of Jane Massey Licata
66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
PAPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
RICHARD APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
APPLICATION NUMBER: 09/048,810
TELECOMMUNICATION NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1454
TELEPHONE: 609-810-1454
                                                                                                                                                                                                                                                                         301 GGAGCAGGCAAATGTGCAAT 320
                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 63, Application US/09280805; Patent No. 6184212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           302 GAGCAGGCAAATGTGCAATA 321
                                                                                                                                                                                                                                                                                               20 GGAGCAGCAAATGTGCAAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GAGCAGGCAAATGTGCAATA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%;
            information for SEQ ID NO:
sequence CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TYPE: TYPE: STRANDEDNESS: Single
ANTI-SERSE: Yes
US-09-280-805-62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
609-810-1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Office
STREET: 66 East Main
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 08053
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-63
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                    RESULT 216
US-09-280-805-63/c
TELEFAX:
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 20;
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Sequence 64, Application US/09280805
Sequence 64, Application US/09280805
Fatent No. 6184212
Fatent No. 6184212
Fatent No. 6184212
Fatent No. 6184210N
Fatent No. 6184210N
Fatent Carbam, Brett P. Monia
FITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
FITLE OF INVENTION: EXPRESSION
FITLE OF SEQUENCES: 271
FORRESPONDENCE ADDRESS: ADDRESS: ADDRESS: ADDRESSE: ADD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WONDPERECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MATCH 26, 1998
ATTOCNEY AGENT INFORMATION:
NAME: Licata, Jame Massey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   32,257
32,257
----- ISPH-0346
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFRENCE/DOCKET NUMBER: IS
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: Nucleic Acid
STRANDEDNESS: Single
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ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         U.S.A.
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Gaps

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ASSULT SEASONS SEASONS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                306 AGGCAAATGTGCAATACCAA 325
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REGISTRATION NUMBER: 32,257
REPRENCE/DOCKET NUMBER: ISPH
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 AGGCAAATGTGCAATACCAA 1
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                 20 base pairs
                                                 TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                             TOPOLOGY: Linear
TI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OPERATING SYSTEM:
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US-09-280-805-66
                     LENGTH:
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APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: NU
COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: U$/09/280,805
FILING DATE: herewith
CLASSIFICATION NUMBER: 09/048,810
FRICR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENI INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 20; DB 1; Li
Pred. No. 3.3e+02;
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COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/48,810
FILING DATE: MARCH 26, 1998
ATTOONEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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REGISTRATION NUMBER: 32,257
REFRENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
INFORMATION FOR SEQ ID NO: 66: SEQUENCE CHARACTERISTICS:
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; Sequence 66, Application US/09280805
; Patent No. 6184212
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 65: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs TYPE: Nucleic Acid STRANDEDNESS: Single TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  / ANTI-SENSE: Yes
US-09-280-805-65
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STATE: NJ
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 Length 20;
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Ouery Match 0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
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US-09-280-805-68/c
; Sequence 68, Application US/09280805
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CURRENT APPLICATION DATA:
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                                                                  CLASSIFICATION:
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; Sequence 69, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
    APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
    APPLICANT: Graham, Brett P. Monia
    TITLE OF INVENTION: EXPRESSION
    NUMBER OF SEQUENCES: 271
    CORRESPONDENCE ADDRESS:
    ADDRESSEE: Law Offices of Jane Massey Licata
    STREET: 66 East Main Street
    CITY: Marlton
    STATE: NJ
    COUNTRY: U.S.A.
    ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
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                GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Grahm, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
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tive 0; Mismatches
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OCHENTE 1484 PC
COMPUTER: 1484 PC
COMPUTER: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY, AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION FOR SEQ ID NO: 68:
SEQUENCE CHARACTERISTICS:
LENGTH 20 Dase pairs
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SOFTWARE: WORDPERFECT 6.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 100.0
Matches 20; Conservative
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STRANDEDNESS: Single
TOPOLOGY: Linear
NTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC OPERATING SYSTEM:
                                                                                                                                                                                                                                                                     U.S.A.
                                                                                                                                                                                                                          CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
  Patent No. 6184212
                                                                                                                                                                                                                                                                                               ZIP: 08053
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APLICATION NUMBER: US09/280,805
FILMS ONTE: herewith 19/089,180
FILMS ONTE: herewith 19/089,180
FILMS ONTE: herewith 19/089,180
FILMS ONTE: herewith 18/09/280,805
FILMS ONTE: herewith 18/09/280
FILMS ONTE: herewith 18
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COUNTRY: U.S.A.
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US-09-280-805-72
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                                                                                                                           Length 20;
                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE Law Offices of Jane Massey Licata STREET: 66 East Main Street CITY: Marlton STATE: No. A. COUNTRY: U.S.A. ZIP: 08053
COUNTRY: U.S.A. ZIP: 08053
CONPUTER READBLE FORM: MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: EARL PC
COMPUTER IN PROPERIES OF COMPUTER: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: BAP PC
COMPUTER: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: BAP PC
COMPUTER: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: BAP PC
COMPUTER: MAPPE
COMPUTER: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE CORRESTED TO NUMBER: US/09/280,805
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licate, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELLECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 72, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
                                                                                                                     Query Match
0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-280-805-71/c
; Sequence 71, Application US/09280805
; Patent No. 6184212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                311 AATGTGCAATACCAACATGT 330
                                                                                                                                                                                                                                          310 AAATGTGCAATACCAACATG 329
                                                                                                                                                                                                                                                                                                  20 AAATGTGCAATACCAACATG 1
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INFORMATION FOR SEQ ID NO: 71:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 100.
Matches 20, Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-280-805-72/c
; TOPOLOGY:
; ANTI-SENSE:
US-09-280-805-70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; ANTI-SENSE:
US-09-280-805-71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
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| Sequence 73, Application US/09280805
| Patent No. 6184212
| GENERAL INCOMMATION:
| APPLICANT: Loraham, Brett P. Monia | TITLE OF INVENTION: EXPRESSION | TITLE OF INVENTION: EXPRESSION | NUMBER OF SEQUENCES: 27 | CORRESPONDENCE ADDRESS: ADDRESSEE: Law Offices of Jane Massey Licata | STREET: 66 East Main Street | CITY: Marlen | STREET: 65 East Main Street | STREET: 67 EAST | STREET: 67 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 20;
                                                   ANTISENSE MODULATION OF HUMAN MDM2 EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: NJ
CONNTRY: U.S.A.

CONPUTRY: U.S.A.

COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
COMPUTE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
RIBING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 20; DB 1; L. Pred. No. 3.38+02;
                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODU
TITLE OF INVENTION: EXPRESSION
TUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane P
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              312 ATGTGCAATACCAACATGTC 331
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REGISTRATION NUMBER: 32,257
REPERENCE/DOCKET NUMBER: ISPH-
TELECOMMUNICATION INFORMATION:
TELEPANE: 609-810-1515
INFORMATION FOR SEQ ID NO: 72:
SEQUENCE CHARACTERISTICS:
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TYPE: Nucleic Acid
STRANDEDNESS: Single
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TOPOLOGY: Linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 229
US-09-280-805-76/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 74, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERPECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
STREET: Marlton
STATE: NJ
              PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTOCNEY, AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REPERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIPICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILIG DATE: March 26,1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECHIONE: 609-810-1515
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| INFORMATION FOR SEQ ID NO: 74: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs | TYPE: Nucleic Acid | STRANDEDNES: Single | TOPOLOGY: Linear | ANTI-SENSE: Yes
                                                                                                                                                                                                                  TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 73:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: SINGle
                                                                                                                                                                                                                                                                                                                                                   , ANTI-SENSE: Yes
US-09-280-805-73
CLASSIFICATION:
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; Sequence 75, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: EXPRESSION
; TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; CITY: Marlton
Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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Patent Wo. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTIENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITATE: MAILEON
STATE: MOSS

ZIP: 08053
COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Lidata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKT NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPAX: 609-810-1515
TELEPAX: 609-810-1515
TELEPAX: 609-810-1515
TELEPAX: 609-810-1515
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Best Local Similarity 100.0%; Pred. No. ...
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     323 CAACATGTCTGTACCTACTG 342
                                                                                                      314 GTGCAATACCAACATGTCTG 333
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0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02;
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Sequence 78, Application US/09280805

Patent No. 6184212

Gardent INFORMATION:
GAPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2;
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:

MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
SOFTWARE: WARDPERFECT 6.0
CURRENT INCHAR: US/09/280,805
FILING DATE: herewith
CLASSIFICATION NUMBER: US/09/280,805
FILING DATE: MATCH 26, 1998
ATTORNEY CAPATION NUMBER: 09/048,810
FILING DATE: MATCH 26, 1998
ATTORNEY CAPATION NUMBER: 32,257
REFERENCE, DOCKET UNUMBER: 12,257
REFERENCE, DOCKET UNUMBER: 32,257
REFERENCE, DOCKET UNUMBER: 12,874
TELEFHONE: 609-810-1515
TELEFHONE: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 609-810-1515
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TELEFAX: 609-810-1515
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0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Law Offices of Jane Massey Licata
STRET: 66 East Main Street
CITY: Marlton
STATE: NJ
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 77:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       351 GTAACCACCTCACAGATICC 370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 GTAACCACCTCACAGATTCC 1
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TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ) TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-77
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: U.S.A.
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APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Breat P. Monia
ATTLE OF INVENTION: EXPRESSION
INTRER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 6 East Main Street
CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
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                                                                                                                                                                                                                                                COUNTA.,
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
MINITER: IBM PC
MINITER: STORAGE
                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOCTHWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIPICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
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; Sequence 77, Application US/09280805
; Patent No. 6184212
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INFORMATION FOR SEQ ID NO: 7
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
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Matches 20; Conservative
        NUMBER OF SEQUENCES: 271
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                                                                                                                                                        CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; TOPOLOGY: Linear; ANTI-SENSE: Yes US-09-280-805-76
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US-09-280-805-80
                                                                 COUNTRY:
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US-09-280-805-79/C
; Sequence 79, Application US/09280805
; Patent No. 6164212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.;
; TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
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  0; Indels
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APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1; Le
. 3.3e+02;
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0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: WORDERFEET 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTONNEY, AGENT INPORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECOMUNICATION INFORMATION:
TELECOMUNICATION NO. 79:
SEQUENCE CHARACTERISTICS:
LENGTH 20 Date pairs
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                                           361 CACAGATTCCAGCTTCGGAA 380
                                                                 20 CACAGATTCCAGCTTCGGAA 1
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20; Conservative
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STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
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Matches
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; Sequence 81. Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
    APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia
    TITLE OF INVENTION: EXPRESSION
    NUMBER OF SEQUENCES: 271
    CORRESPONDENCE ADDRESS:
    ADDRESSEE: Law Offices of Jane Massey Licata
    STREET: 66 East Main Street
                                                                                COMPUTER READABLE FORM:

MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC
COPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDERFECT 6.0
CURRENT PPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY,AGRAY INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE,DOCKET NUMBER: 15PH-0346
TELECOMMUNICATION INPORMATION:
TELEBRING: 609-810-1115
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ZIP: 08053
COMPUTER READBLE FORM:
MEDIN TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IEM PC
OPERATING SYSTEM: WINDOWS 95
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Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
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herewith
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APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY AGENT INFORMATION:
NAME: Licata, Jane Massey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    386 GACCCTGGTTAGACCAAAGC 405
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66 East Main Street
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 8
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: Linear ANTI-SENSE: Yes
                                                                      U.S.A.
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                      Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Marlton
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STATE: NJ
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ZIP: 08053
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US-09-280-805-84/c
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Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                          0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 82, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRIES
ZIP: 08053
COMPOTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
CAMPITER: IBM PC
WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3a+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street STREET: Marlton STATE: NJ
                                                                                                                                                                                                                                                                                             Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.: Matches 20; Conservative 0; Mismatches
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION: INFORMATION:
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 81:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ISPH-0346
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OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERECT 6.0
SUPTRAIN APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-034
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
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SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
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US-09-280-805-82
                                                                                                                                                                                                               ; TOPOLOGY: Linear; ANTI-SENSE: Yes US-09-280-805-81
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US-09-280-805-83/c
; Sequence 83, Application US/09280805
; Patent No. 6184212
; Patent No. 6184212
; Patent No. 6184212
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
; TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: LAW OFFICES of Jane Massey Licata
                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC OPERATING SYSTEM: WINDOWS 95
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66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION WARSE: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
RAGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; Sequence 84, Application US/09280805; Patent No. 6184212
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TELEPHONE: 609-810-1515
TELEFAX: 609-810-1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO: 8
SEQUENCE CHARACTERISTICS:
TENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
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609-810-1515
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  TELEPHONE:
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US-09-280-805-86
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Locen J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERRECT 6.0
SOFTWARE: WORDPERRECT 7
APPLICATION NUMBER: US/09/280,805
PILING DATE:
                 ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match . 0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                              CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
PLING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licea, Jane Massey
REGISTRATION NUMBER: 32 257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECHONE: 609-810-1515
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
                                                                                                                                                           APPLICATION NUMBER: US/09/280,805 FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       450 ACTTATACTATGAAAGAGGT 469
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-280-805-85/c; Sequence 85, Application US/09280805 Patent No. 6184212; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 84: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs TYPE: Nucleic Acid STRANDEDNESS: Single TOPOLOGY: Linear ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 ACTTATACTATGAAGAGGT 1
                                                                          COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CLASSIFICATION:
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-280-805-84
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                                                                                                                                                                                                                                                             Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 86, Application US/09280805
Fatent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: BYPRESSION
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCES 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                                                                                                                                                                                                                                                       Query Match 0.8%; Score 20; DB 1; L
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Logal Similarity 100.0%; Pred. No. 3.3
Matches 20, Conservative 0, Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION
APPLICATION
APPLICATION
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                        477 TATCTTGGCCAGTATATTAT 496
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     490 ATATTATGACTAAACGATTA 509
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISJ
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 8
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 20 base par
TYPE: Nucleic Acid
                                                                                                                                                    TOPOLOGY: Linear
HANTI-SENSE: Yes
US-09-280-805-85
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GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Coren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brette P. Monia
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1; Le
3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                          SCPTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
PPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTONNEY AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDFERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MATCH 26,1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ISPH-0346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 89, Application US/09280805
Patent No. 6184212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     503 ACGATTATATGAGAGG 522
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 ACGATTATATGATGAGGG 1
                      COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,2
                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 609-810-1454
TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      , ANTI-SENSE: Yes
US-09-280-805-88
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         U.S.A.
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US-09-280-805-89/C
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ZIP: 08053
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Sequence 88, Application US/09280805

Sequence 88, Application US/09280805

GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Breett P. Monia
TITLE OF INVENTION: EXPRESSION
TITLE OF INVENTION: EXPRESSION
CORRESPONDENCE: 271
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 20;
                                                     Sequence 87, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Craham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESSED LAW OFFICES OF Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 20; DB 1; Le
Pred. No. 3.3e+02;
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0.8%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPREFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFRENCE/DOCKET NUMBER: ISPH
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPAX: 609-810-1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20 reacraaacearrararear 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
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CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                    CITY: Marlton
STATE: NJ
                                       US-09-280-805-87/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-280-805-87
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Length 20;

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Marlton
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US-09-280-805-92/c
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                                                                                                                                                                                                                                                                 KESULT 243
US-09-280-805-90/C
Sequence 90, Application US/09280805
PATENT NO. 6184212
GRNERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
CONTEST NA
                                                                                                                                                              Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                DB 1; Le
                                                                                                                                                            Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IEM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERPECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRICK APPLICATION DATA:
APPLICATION UNMBER: 09/048,810
FILING DATE: MARCH 26,1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                           515 TGAGAAGCAACAACATATTG 534
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            525 CAACATATTGTATATTGTTC 544
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INFORMATION FOR SEQ ID NO: 90:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
; SEQUENCE CHARACTERISTICS:
; LENGTH: 20 base pairs
; TYPE: Nucleic Acid
; STRANDEDNESS: Single
; ANTI-SENSE: Yes
US-09-280-805-89
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-90
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RESULT 244 US-09-280-805-91/c

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Gaps
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; Sequence 92, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2; TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 27; CORRESPONDENCE ADDRESSS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
; CITY: Marlton
; STATE: 'N J.
; COUNTRY: U.S.A.
ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02;
Sequence 91, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Breet P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                              MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Prec. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   531 ATTGTATATTGTTCAAATGA 550
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REPERENCE/DOCKET NUMBER: 1SPH-
TELECOMMUNICATION INPORMATION:
TELEPAN: 609-810-1155
INFORMATION FOR SEQ ID NO: 91:
SEQUENCE CHARACTERISTICS:
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Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 20 Dasc TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                          COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: Linear
HANTI-SENSE: Yes
US-09-280-805-91
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RESULT 248
US-09-280-805-95/c
; Sequence 95, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                         STRANDEDNESS: Single
                                                                   Linear
Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Linear
Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             U.S.A.
                                                               ; TOPOLOGY:
; ANTI-SENSE:
US-09-280-805-93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ANTI-SENSE:
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                                                                                                                                                                                                                                                                                                                                            RESULT 247
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Patent No. 6184212

GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
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ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
CAMPITER: IBM PC
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey,
REFERENCE/DOCKET NUMBER: 12,257
REFERENCE/DOCKET NUMBER: ISPH-0346
FILEPHONE 609-810-1515
TELEPHONE 609-810-1515
TYPE: NUCleic Acid
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
108-09-280-805-92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1454
INFORMATION FOR SEQ ID NO: 93:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
PRICR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     538 ATTGTTCAAATGATCTTCTA 557
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 ATTGTTCAAATGATCTTCTA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.81
Best Local Similarity 100.0
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-280-805-93/c
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                                                           Gaps
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     Length 20;
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                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STARES: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYER READABLE FORM:
MEDIUM TYER: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MATCh 26, 1998
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION SOFT STORES SOFT SECURITIES:
SEQUENCE CHARACTERISTICS:
TENTALLY 20 DASE DAILE
     Score 20; DB 1; Le
Pred. No. 3.3e+02;
Query Match

0.8%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
                                                                                                          549 GATCTTCTAGGAGATTTGTT 568
                                                                                                                                                                                                                                                                     US-09-280-805-94/c
; Sequence 94, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
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Sequence 97, Application US/09280805; Patent No. 6184212; GENERAL INFORMATION: GENERAL INFORMATION: APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia TITLE OF INVENTION: AATISENSE MODULATION OF HUMAN MDM2 TITLE OF INVENTION: EXPRESSION NUMBER OF SEQUENCES: 27 CORRESPONDENCE ADDRESS: ADDRESSE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
       TLILING DAILS:

CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INPORMATION:
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1515
TELEPHONE: 20 Dase pairs
TELENTY: 20 Dase pairs
TENNOTH: 20 Dase pairs
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26,1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               575 GCCAAGCTTCTCTGTGAAAG 594
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OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 9
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
  FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: Linear;
ANTI-SENSE: Yes
US-09-280-805-96
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: NJ
COUNTRY: U.
ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-280-805-97/c
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100.0%; Pred. No. 3.3e+02;
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Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Araham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLED OF INVENTION: EXPRESSION
                     APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: NATION.
STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                                                                                                                                                                                                                                                                      ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
Loren J. Miraglia, Pamela Nero, Mark J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 Bast Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
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SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
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TELBEAK: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
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TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: Linear
TI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-280-805-96/c
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US-09-280-805-95
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PRIOR APPLICATION DATA:
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Sequence 98, Application US/09280805
Fatent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Mailton
                                          0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 20; DB 1; Length 20;
Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 99, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  100.0%; Pred. ...
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COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                           587 TGTGAAAGAGCACAGGAAAA 606
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             593 AGAGCACAGGAAAATATA 612
                                                                                                                                                                   20 TGTGAAAGAGCACAGGAAAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 AGAGCACAGGAAAATATA 1
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INFORMATION FOR SEQ ID NO: 9
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                     Query Match 0.8°
Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 20; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Linear
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US-09-280-805-98
US-09-280-805-97
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
                                                                                                                                                         DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                            ADDRESSEB: Law Offices of Jane Massey Licata STREET: 66 East Main Street CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                MEDIUM TIPE: DISKETIE, 3.5 INCH, 1.44
COMPUTER: 1BM PC
OPERATING SYSTEM: WINDOWS 95
SOSTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION.
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTONNEY, AGENT INPORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECHONE: 609-810-115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: WORDPERPECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  600 AGGAAAATATATACCATGAT 619
EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 AGGAAATATATACCATGAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: 99: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs
              NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TOPOLOGY: Linear; ANTI-SENSE: Yes
US-09-280-805-99
                                                                                                      COUNTRY: U.S.A.
ZIP: 08053
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 101, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: NO. COUNTRY: NO. COUNTRY: NO. COUNTRY: NO. S.A. COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC OPERATING SYSTEM: WINDOWS 95 SOFTWARE: WORDPERFECT 6.0 CURRENT APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: PRIOR APPLICATION DATA: PRIOR APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION DATA: APPLICATION NUMBER: 09/048,810 FILING DATE: MATCH 26, 1998 ATTORNEY/AGENT INFORMATION:
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ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                    NAME: Licata, Jame Massey
REGISTRATION NUMBER: 32,257
REFRENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
INFORMATION FOR SEQ 1D NO: 100: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ISPH-0346
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             609 TATACCATGATCTACAGGAA 628
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REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 151
TELECOMMUNICATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 101:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Nucleic Acid
EDNESS: Single
                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: Linear; ANTI-SENSE: Yes
US-09-280-805-100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: Marlton
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DB 1; Length 20;

0.8%; Score 20;

Query Match

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GENERAL INFORMATION:

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.

APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: ANTIGENSE MODULATION OF HUMAN MDM2

TITLE OF INVENTION: EXPRESSION

NUMBER OF SEQUENCES: 271

CORRESPONDENCE ADDRESS:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street

CITY: Marlton

STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                       Indels
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Sequence 103, Application US/09280805
Sequence 103, Application US/09280805
Sequence 103, Application US/09280805
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READBLE FORM:
MEDIUTER: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: ISM PC
OOMPUTER: ISM PC
OPERATING SYSTEM: WINDOWS 95
100.0%; Pred. No. 3.3e+02;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAMME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              REFERENCE/DOCKET NUMBER: ISPH-0346
                                                                                                                                                                                                                                              ; Sequence 102, Application US/09280805; Patent No. 6184212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          634 TAGTAGTCAATCAGCAGGAA 653
                                                                                                           20 TCTACAGGAACTTGGTAGTA 1
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                                                                       619 TCTACAGGAACTTGGTAGTA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTER.STICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TOPOLOGY: Linear; ANTI-SENSE: Yes
US-09-280-805-102
                                                                                                                                                                                             RESULT 255
US-09-280-805-102/c
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Query Match
0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                    Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REPERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1554
INFORMATION FOR SEQ 1D NO: 104:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
INFORMATION FOR SEQ 1D NO: 1CH:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.4
COMPUTER: IBM PC
OMPUTER: IBM PC
OPPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION: Herewith
PRIOR APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
RESISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
                                                                                                                                                                                                                                                                                                                                                                                                                                                 656 ATCGGACTCAGGTACATCTG 675
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 Arcedacreageracarere 1
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                           TOPOLOGY: Linear
HANTI-SENSE: Yes
US-09-280-805-104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               U.S.A.
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ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-280-805-105
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CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
WEDTUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FTIING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 104, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION
APPLICANT: Coren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2 TITLE OF INVENTION: EXPRESSION CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
                                                                                                                ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Pred. No. 3.3e+02;
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                          CURPLAINS

CURPLAINS SYSTEM: WINDOWS >> SOFTWARE: WORDERFRET 6.0

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/280,805

FILING DATE: herewith

CLASSIFICATION:

PRIOR APPLICATION NUMBER: 09/448,810

FILING DATE: March 26, 1998

APPLICATION NUMBER: 32,257

REGISTRATION NUMBER: 32,257

REGISTRATION NUMBER: 32,257

REFERENCE/DOCKET NUMBER: 1SPH-0346

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION 103:

SEQUENCE CHARACTERISTICS:

LENGTH: 20 base pairs

TYPE: Nucleic Acid

STRANDEDDENS: Single

TOPOLOGY: Linear

105-09-280-805-103
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
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Best Local Similarity 100.0°
Matches 20; Conservative
                                                                                           U.S.A.
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                                                                                        COUNTRY: U
ZIP: 08053
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                                                                     STATE:
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                                                                                                                                                                                                 Sequence 105, Application US/09280805
Fatent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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 DB 1; Le
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Length 20;
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APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata
STREET: Mariton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPTRY: UOS.A.

ZIP: 08053
COMPUTER READDBLE PORM:
MEDIUM TYEE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION DATA:
RICHARD AND STATES AND STATE
                                                                                                                                                        MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Le
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0.8%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                  SOFTWARE: WORDPERFECT 6.0 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/48,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECHONE: 609-810-1515
TELECHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 261
US-09-280-805-108/c
; Sequence 108, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              691 GTCACCTTGAAGGTGGGAGT 710
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 107: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs TYPE: Nucleic Acid STRANDEDNESS: Single
                                                                                                                  COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: Linear
ANTI-SENSE: Yes
                                      U.S.A.
                                                                        08053
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                                          COUNTRY:
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                                                                                                                                                                                                                                                          Sequence 106, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTAL.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CONTIDENT INTO THE STATE OF THE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ISPH-0346
669 ACATCTGTGAGTGAGACAG 688
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TELECOMMUNICATION INFORMATION:
TELEFONE: 609-810-1515
TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 106:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 AGAACAGGTGTCACCTTGAA 1
                                                  20 ACATCTGTGAGTGAGAACAG 1
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Best Local Similarity 100.0
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Marlton
                                                                                                                                                                                                                                         US-09-280-805-106/c
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COUNTRY:
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S-09-280-805-110/c
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COUNTRY: U.
ZIP: 08053
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 109-209/C
Sequence 109-209/C
Patent No. 6184212
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CONTY: MAILON
CONTY: MAILON
                                                                                                                                                                                                                                              Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Pred. ....
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APPLICATION NUMBER: 0/048,810
FILING DATE: March 26, 1998
ATTORNEY/ACENT INFORMATION:
NAWE: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 15PH-034
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                             704 TGGGAGTGATCAAAAGGACC 723
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1454
INFORMATION FOR SEQ ID NO: 108:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
TOPOLOGY: Linear
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
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TYPE: Nucleic Acid
STRANDEDNESS: Single
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; ANTI-SENSE: Yes
US-09-280-805-109
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Best Local Similarity
Matches 20; Conserv
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                                                                                                                                                                                                        US-09-280-805-108
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; Sequence 111, Application US/09280805
; Patent No. 6184212
; Patent No. 6184212
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
; TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
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                                                                            APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION: DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFRONE: 609-810-1515
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3.3e+02;
                                                                                                                                                                                                                                                ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match

0.8%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
Sequence 110, Application US/09280805 Patent No. 6184212 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               727 TACAAGAGCTTCAGGAAGAG 746
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SEQUENCE CHRACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-110
                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A. ZIP: 08053
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                                                                                                                                                                                                               Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 Bast Main Street
CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC
OPPRATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: WORDDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERNICE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-280-805-113/c
; Sequence 113, Application US/09280805
; Patent No. 6184212
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            112:
  ; INFORMATION FOR SEQ ID NO: 1
; SEQUENCE CHARACTERISTICS:
; LENGTH: 20 base pairs
; TYPE: Nucleic Acid
; TYPE: Nucleic Acid
; STRANDEDNESS: Single
; ATOPOLOGY: Linear
; ATTI-SENSE: Yes
US-09-280-805-112
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 20 base pairs
TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: Linear
HANTI-SENSE: Yes
US-09-280-805-113
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; Sequence 112, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: EXPRESSION
; TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; CITY: Marlton
; CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.38+02;
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 08053

COMPUTER READBLE FORM:
MEDIUM TYPE. DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        100.0%; Pred. ...
                                             COMPUTER: 1BM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARES WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MATCH 26,1998
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          740 GGAAGAAACCTTCATCTT 759
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ATTORNEY/AGENT INCOMESSEY
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPAX: 609-810-1514
INFORMATION FOR SEQ ID NO: 111:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 GGAAGAGAACCTTCATCTT 1
                                                                                                                                                                                                                                                                                                           NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.8 Best Local Similarity 100. Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   609-810-1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; ANTI-SENSE: Yes
US-09-280-805-111
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: NJ
COUNTRY: U.S
ZIP: 08053
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US-09-280-805-116/c
; Sequence 116, Application US/09280805
Patent No. 6184312
; GENERAL INFORMATION:
                                                                                FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     787 CATCTAGAAGGAGAGCAATT 806
                                                                                                                                                                                                   NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-
TELECOMMUNICATION INFORMATION:
TELEPAN: 609-810-1515
INFORMATION FOR SEQ ID NO: 115:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match

Best Local Similarity 100.0%; P

Matches 20; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 CATCTAGAAGGAGGAGTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 116:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                 ; LENGTH: 20 base pairs
; TYPE: Nucleic Acid
; STRANDEDNESS: Single
; TOPOLGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 269
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                                                   GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
APPLICANT: Graham, Brett P. Monia
APPLICANT: Graham, ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Malton
STATE: N. COUNTRY: U.S.A.
COUNTRY: U.S.A.
STATE: DISKETTE: ADSRESTED SEGUENCES
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             100.0%; Preu. ...
                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC
COMPUTER: OF CONTROL SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION NUMBER: 09/048,310
PRIOR APPLICATION NUMBER: 09/048,310
PRIOR APPLICATION NUMBER: 126,1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 13.257
REFERENCE/DOCKET NUMBER: 13PH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
INFORMATION FOR SEQ ID NO: 114:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
LENGTH: 20 base pairs
TELENGTH: 20 base pairs
TERMENDENESS: Single
                    Sequence 114, Application US/09280805
Patent No. 6184212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               774 AGACCATCTACCTCATCTAG 793
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20 AGACCATCTACCTCATCTAG 1
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Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; TOPOLOGY: Linear; ANTI-SENSE: Yes US-09-280-805-114
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TRY: U.S.A.
08053
US-09-280-805-114/c
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Length 20;
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: NO COUNTRY: NO S.A.

ZIP: 08053

COMPUTER READBALE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
OPREATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Herewith
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MATCH 26, 1998
ATTORIEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 20; DB 1; L
100.0%; Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
STREET: Narlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                              ISPH-0346
SUCKENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
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SENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                               | Sequence 117, Application US/09280805
| Patent No. 6184212
| GENERAL INFORMATION:
| APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
| APPLICANT: Graham, Brett P. Monia
| TITLE OF INVENTION: EXPRESSION
| TITLE OF INVENTION: EXPRESSION
| TITLE OF INVENTION: EXPRESSION
| TITLE OF SEQUENCES: 271
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: Law Offices of Jane Massey Licata
| STREET: 66 East Main Street
| CITY: Mariton
| STREET: Action Main Street
                                                                                                                                    Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U.S....

ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAMM: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPRONE: 609-810-1515
TELEPAX: 609-810-1555
TELEFAX: 609-810-1555
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 271
US-09-280-805-118/c
; Sequence 118, Application US/09280805
; Patent No. 6184212
                                                                                                                                                                                                                              798 AGAGCAATTAGTGAGACAGA 817
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                810 GAGACAGAAGAAATTCAGA 829
                                                                                                                                                                                                                                                                       20 AGAGCAATTAGTGAGACAGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 GAGACAGAAGAAATTCAGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
; TYPE: Nucleic Acid
STRANDEDNESS: Single
; TOPOLOGY: Linear
; ANTL SENSE: Yes
US-09-280-805-116
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: Linear
TI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                     US-09-280-805-117/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; ANTI-SENSE:
US-09-280-805-117
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Gaps
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APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 119, Application US/09280805
Patent No. 6184312
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLER OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: NO
COMPUTER: NO
COMPUTER: NO
COMPUTER READALE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
COMPUTER: NOTE: 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
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                                                                                                                                                                                                                                                ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E: Law Offices of Jane Massey Licata
66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          824 TTCAGATGAATTATCTGGTG 843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 rrcagargaarrarcregre 1
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Best Local Similarity 100.0
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              609-810-1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TOPOLOGY: Linear; ANTI-SENSE: Yes US-09-280-805-118
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-280-805-119/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE:
STREET: 66
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CLASSIPCATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-034
TELEPHONE: 609-810-1515
INFORMATION: FOR SEQ ID NO: 121:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  857 ACGCCACAAATCTGATAGTA 876
                                                                                                                                                  844 AACGACAAAGAAAACGCCAC 863
                                                                                                                                                                                20 AACGACAAAGAAAACGCCAC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        20 ACCCCACAAATCTGATAGTA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            U.S.A.
                                                                                                                                                                                                                                                    RESULT 274
US-09-280-805-121/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-280-805-122/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 08053
      ; ANTI-SENSE:
US-09-280-805-120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    , ANTI-SENSE:
US-09-280-805-121
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 273
US-09-280-805-120/C
i Sequence 120, Application US/09280805
i Sequence 120, Application US/09280805
i Sequence 120, Application US/09280805
i GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
ITILE OF INVENTION:
ITILE OF INVENTION: EXPRESSION
ITILE OF INVENTION: EXPRESSION
INUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
COUNTRY: U.S. R.
ZIP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A.

ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPEE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Liceta, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 15PH-0346
TELECOMMUNICATION INPORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                 MAME: Licata, Jane Massey

REGISTRATION NUMBER: 32,257

REFERENCE/DOCKET NUMBER: 1SPH-0346

TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515

INFORMATION FOR SEQ ID NO: 119:
SEQUENCE CHARACTER FILES

LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear

ANTI-SENSE: Yes
US-09-280-805-119
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                     FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.8%;
Best Local Similarity 100.0%
Matches 20; Conservative
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
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                                                                                                                                                                                                                                                                              Sequence 121, Application US/09280805
| Patent No. 6184212
| GENERAL INFORMATION:
| APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2: TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2: CORRESPONDENCE ADDRESS:
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: Law Offices of Jane Massey Licata | CITY: Mailton
Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER CRADABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOPTWARE: WORDPERFECT 6.0
CARRENT APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%; Score 20; DB 1; Ler
... Pred. No. 3.3e+02;
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0.8%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
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Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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Patent No. 6184212
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS: ADDRE
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ZIP: 08053

COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
                              APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REPERNICE/DOCKET NUMBER: 1SPH-0346
TELEPRONE: 609-810-1515
TELEPRONE: 609-810-1515
INFORMATION FOR SEQ ID NO: 123: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDENESS: Single
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APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                880 CCCTTTCCTTTGATGAAGC 899
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TELEPHONE: 609-810-1515
TELEFAX: 609-810-1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,2
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs TYPE: Nucleic Acid STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 t TOPOLOGY: Linear
t ANTI-SENSE: Yes
US-09-280-805-123
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-280-805-124/c
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Sequence 123, Application US/09280805

Patent No. 6194212

Sequence 123, Application US/09280805

Sequence 123, Application:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: BRET P. Monia
TITLE OF INVENTION: BRET P. Monia
TITLE OF INVENTION: BRET P. Monia
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053

COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPPERATIOS SYSTEM: WINDOWS 95
SOFTWARE: WORDDERFECT 6.0
CURRENT APPLICATION DATA:
STILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
ANTISENSE MODULATION OF HUMAN MDM2 EXPRESSION
                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
                                                            NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM DC
OPERATING SYSTEM: WINDOWS 95
SOSTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION:
APPLICATION:
FILING DATE: WARCH 26, 1998
APPLICATION NUMBER: 09/048,810
FILING DATE: WARCH 26, 1998
ATTORNEY AGENT INFORWATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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INFORMATION FOR SEQ ID NO: 122:
SEQUENCE CHARACTER.STICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             867 TCTGATAGTATT
TITLE OF INVENTION:
TITLE OF INVENTION:
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US-09-280-805-122
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CORRESPONDENCE ADDRESS:
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US-09-280-805-126
                                                                                                                                                                                                              COUNTRY:
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| Sequence 125, Application US/09280805
| Patent No. 6184212
| GENERAL INFORMATION:
| GENERAL INFORMATION:
| APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia
| TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2 |
| TITLE OF INVENTION: EXPRESSION |
| NUMBER OF SEQUENCES: 271 |
| CORRESPONDENCE ADDRESS: | ADDRESSE: Law Offices of Jane Massey Licata |
| STREET: 66 East Main Street |
| CITY: Mailton |
| COTY: Mailton |

Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.38+02; Matches 20; Conservative 0; Mismatches 0; Indels
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US-00-280-055-126/c
Sequence 126, Application US/09280805
Sequence 126, Application US/09280805
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Breet P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2:
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE:
COMPUTER: IBM 90053
COMPUTER: IBM 90058
SOFTWARE: WORDPREFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: March 100.
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                895 AAAGCCTGGCTCTGTGTA 914
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INFORMATION FOR SEQ ID NO: 125:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 TOPOLOGY: Linear
1 ANTI-SENSE: Yes
US-09-280-805-125
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COUNTRY:
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; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2 TITLE OF INVENTION: EXPRESSION
; VINDER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
; CITY: Marlton
; STATE: NJ
; COUNTRY: U.S.A.
; COMPUTER READABLE FORM:
; MARDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
                                                                                                                                    COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION:
PRIOR APPLICATION:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INPORMATION:
NAME: Licata, Jame Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REJERBNCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1155
TELEPHONE: 609-810-1156
ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Preu. ...
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herewith
J.
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APPLICATION NUMBER: 09/048,
FILING DATE: March 26, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.C
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: Linear
TI-SENSE: Yes
                                                                                                  U.S.A.
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; Sequence 130, Application US/09280805
; Patent No. 6184212
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: ANTIERNE MODULATION OF HUMAN MDM2
; TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.38+02;
                                                                                                                                                                                    Sequence 129, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Craham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2;
TITLE OF INVENTION: APTISENSE MODULATION OF HUMAN MDM2;
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREST: 66 Bast Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U.S.A.

ZIP: 08053
COMPUTER READALE FORM:
MODIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                         936 AGAAGCAGTAGCAGTGAATC 955
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             949 GTGAATCTACAGGGACGCCA 968
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TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 GTGAATCTACAGGGACGCCA
                                                                   20 AGAAGCAGTAGCAGTGAATC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,2
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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Matches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Marlton
                                                                                                                                                                  US-09-280-805-129/c
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; Sequence 128, Application US/09280805
; Patent No. 6184212
; Patent No. 6184212
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
; TITLE OF INVENTION: EXPRESSION
; CORRESPONDENCE ADDRESS:
; CARRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
; CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                     0; Indels
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MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                       Score 20; DB 1; Le
Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CUMPUTER: 14M PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CUMRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTONNEY, AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                      ISPH-0346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  927 TGTTGTGAAAGAAGCAGTAG 946
                    NAME: Licata, Jane Massey
REGISTRATION UNUMBER: 32,257-
REFERENCE/DOCKET NUMBER: ISPH-
TELECOMMUNICATION INFORMATION:
TELEFHONE: 609-810-1515
TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 127:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 TGTTGTGAAGAAGCAGTAG 1
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ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              609-810-1454
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STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                   TOPOLOGY: Linear
ANTI-SENSE: Yes
US-09-280-805-127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: Linear
TI-SENSE: Yes
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TELEFAX: 6
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US-09-280-805-128
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Gaps

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Sequence 132, Application US/09280805
Fatent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Mariton
                                                                                                                                                                                                                                                                                                                                  Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: 1BM PC
OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.8%; Score 20; DB 1; L
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
  REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: WORDPERPECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                     971 GAATCCGGATCTTGATGCTG 990
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 GAATCCGGATCTTGATGCTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISE
TELECOMMUNICATION:
TELEPHONE: 609-810-1515
                                                                         TELEFAX: 609-810-1454
INPORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNES: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE: 609-810-151
TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TOPOLOGY: Linear; ANTI-SENSE: Yes
US-09-280-805-132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U.S.A. ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     JS-09-280-805-132/c
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US-09-280-805-131
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                                                            COMPUTER READABLE FORM:

MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: DISKETTE, 3.0

CURREAL: WORDPERFECT 6.0

CURREAL: APPLICATION NUMBER: US/09/280,805

FILING DATE: herewith
CLASSIFICATION NUMBER: 09/048,810

FILING DATE: MARCH 26, 1998

ATTORNEY, AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257

REFERENCE/COCKET NUMBER: 32,257

REFERENCE/COCKET NUMBER: 15.PH-0346

TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1155

TELECOMMUNICATION INFORMATION:
TELEPRAN: 609-810-1154

INFORMATION FOR SEQ ID NO: 130: SEQUENCE CHARACTERISTICS:
LIGHGTH: 20 base pairs
TYPE: Nucleic Acid
STANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC CORENT APPLICATION NUMBER: US/09/280,805

CURRENT APPLICATION NUMBER: US/09/280,805

FILING DATE: herewith CLASSIFICATION NUMBER: 09/048,810

PRIOR APPLICATION NUMBER: 09/048,810

PILING DATE: March 26, 1998

ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Masseey

REGISTRATION NUMBER: 32,257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match

0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street CITY: Marlton STATE: N. STATE: N. U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 284
US-09-280-805-131/c
; Sequence 131, Application US/09280805
; Patent No. 6184212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  964 CGCCATCGAATCCGGATCTT 983
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 CGCCATCGAATCCGGATCTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: Linear
ANTI-SENSE: Yes
                                               U.S.A.
Marlton
                                            COUNTRY: U
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-280-805-130
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Gaps .; 0 983 TGATGCTGGTGTAAGTGAAC 1002

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Sequence 115, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
; TITLE OF INVENTION: 27PRESSION
; UNMBER OF SEQUENCES: 27
; CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
; CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION NUMBER: US/09/48,810
FILING DATE: March 26, 1998
ATTORING APPLICATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                             MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC OPERATING SYSTEM: WINDOWS 95
                                                                                                                  GPERATING SIGLEM: WINDOWS 32 SOFTWARE: WORDPERFECT 6.0 CURRENT APPLICATION DATA: APPLICATION DATA: APPLICATION NUMBER: US/09/280,805 FILING DATE: herewith CLASSIFICATION: PRIOR APPLICATION UNDER: 09/048,810 FILING DATE: MARCH 26,1998 ATTORNEY/AGENT INFORMATION: NAME: Licata, Jane Massey REGISTRATION NUMBER: 32,257 REFENCE/DOCKET NUMBER: 1SPH-0346 TELECOMMUNICATION: TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                 ISPH-0346
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SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 100.
Matches 20; Conservative
                             COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: Linear
HANTI-SENSE: Yes
US-09-280-805-134
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COUNTRY: U.
ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       요
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; Sequence 134, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; GENERAL INFORMATION:
; APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: EXPRESSION
; TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; CITY: Marlton Main Street
; CITY: Marlton Marian Street
; CITY: Marlton Marian Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                               APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Herewith
CLASSIFICATION: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MATCH 26, 1998
ATTORNEY,AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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                                                                                            US-09-280-805-133/c
; Sequence 133, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 133:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 AGTGAACATTCAGGTGATTG 1
20 TGATGCTGGTGTAAGTGAAC
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NJ
... U.S.A.
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ANTI-SENSE: Yes
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COUNTRY: U.S.A.
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Query Match
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                                                                                                                                                                                             DB 1; Length 20; 3.3e+02;
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| Sequence | 136, Application US/09280805
| Patent No. 6184212
| GENERAL INFORMATION:
| APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia
| TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2 TITLE OF INVENTION: EXPRESSION
| NUMBER OF SEQUENCES: 271
| CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
COMPUTER: TAM PC
COMPUTER: MORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFTATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MATCH 26, 1998
ATTORNEY/AGENT INFORMATION:
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.00.0%; Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: 66 East Main Street
CITY: Mariton
STATE: NJ
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                                                                                                                                                                                             Query Match

0.8%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
INFORMATION FOR SEQ ID NO: 136:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                             1017 TTGGATCAGGATTCAGTTTC 1036
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANBEDNESS: Single
TOPOLOGY: Linear
MATI-SENSE: Yes
US-09-280-805-135

    TOPOLOGY: Linear
    ANTI-SENSE: Yes
US-09-280-805-136

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100.0%; Pred. No. 3.3e+02;
Sequence 137, Application US/09280805
Fatent No. 6184212
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
ITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 138, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION:
TITLE OF INVENTION: EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                          DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                  ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
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STREET: 66 East Main Street
CITY: Marlton
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CURRENT APPLICATION DATA:
APPLICATION DATA:
CLASSIFICATION DATA:
PRIOR APPLICATION:
PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
ATTONNEY, AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION SEG 10-1454
INFORMATION FOR SEG 10 NO: 137:
SEQUENCE CHARACTERISTICS:
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Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Office
                                                                                                                                                                                                                                                                                                                                                                                                            IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                    OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                 U.S.A.
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                                                                                                                                                                                                                                                                     CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 291
US-09-280-805-138/c
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US-09-280-805-140/C
; Sequence 140, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; PAPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; TITLE OF INVENTION: EXPRESSION
; TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESSE:
; ADDRESSES: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
; CITY: Marlton
                                                                                                                                                           0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SOFTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Herewith
CLASSIFICATION TO THE PRIOR PELICATION TO THE PRIOR PELICATION TO THE PARTORNEY OF 6, 1998
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                        1051 TAGAATTTGAAGTTGAATCT 1070
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; LENGTH: 20 base pairs
; TYPE: Nucleic Acid
; STRANDEDNESS: Single
; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-139
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                          20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                Best Local Similarity
Matches 20; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: Linear ANTI-SENSE: Yes
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STATE:
                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                        RESULT 293
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 139, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: U.S.A.
ZIP: 08053.
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDFREFCT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION WHERE: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 15SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION SEQ 1D NO: 138:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: NUCLEIC Acid
TYPE: NUCLEIC Acid
TYPE: STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1046 TAGTGTAGAATTTGAAGTTG 1065
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INPORMATION:
                   SYSTEM: WINDOWS 95
WORDPERFECT 6.0
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COMPUTER: IBM PC
OPERATING SYSTEM:
SOFTWARE: WORDPER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TOPOLOGY: Linear; ANTI-SENSE: Yes US-09-280-805-138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
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ö Gaps ö Query Match

0.8%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0; Indels RESULT 294 US-09-280-805-141/c ; Sequence 141, Application US/09280805

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US-09-280-805-143/c
; Sequence 143, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; TITLE OF INVENTION:
; TITLE OF INVENTION: EXPRESSION
; TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
   ADDRESSES: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
; CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
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MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFTCATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 15PH-0346
TELECOMMUNICATION NUMBER: 15PH-0346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                PRIOR APPLICATION DATA:

APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGRAT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 32,257
REPERENCE/DOCKET NUMBER: 1SPH-0346
FILECOMMUNICATION INFORMATION:
FILEPRAM: 609-810-1515
FILEFRAM: 609-810-1515
FILEMATION FOR SEQ ID NO: 142:
BENCHARACTERISTICS:
LENGTH: 20 base pairs
FILENGTH: 20 base pairs
FIREMEN: 20 base pairs
FIREME
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1077 TCAGAAGATTATAGCCTTAG 1096
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTER.STICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                U.S.A.
                                                                                                             CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U
ZIP: 08053
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i Sequence 142 Application US/09280805

i Sequence 142 Application US/09280805

i Sequence 142 Application

i Sequence 142 Application

i Sequence 142 Application

i APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.

APPLICANT: Graham, Brett P. Monia

ITILE OF INVENTION: BYPRESSION

NUMBER OF SEQUENCES:

I TILE OF INVENTION: EXPRESSION

NUMBER OF SEQUENCES:

ADDRESSE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street

CITY: Marlton

STATE: NJ

COUNTRY: U.S.A.

IPP: 08053

COMPUTER READABLE FORM:

MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE

COMPUTER: IBM PC

COMPUTER: NORDPERFECT 6.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
               GENERAL INFORWATION:

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.

APPLICANT: Graham, Breet P. Monia

TITLE OF INVENTION: EXPRESSION

TITLE OF INVENTION: EXPRESSION

NUMBER OF SEQUENCES: 271

CORRESPONDENCE ADDRESS:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street

CITY: Mailton
                                                                                                                                                                                                                                                                                                                                        CITY: Marico..
STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
INFORMATION FOR SEQ ID NO: 141:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TYPE: Linear
ANTI-SENSE: Yes
US-09-280-805-141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IEM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
WANTER FACEL AND ATTORNEY AND ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1068 TCTCTCGACTCAGAAGATTA 1087
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0
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; Patent No. 6184212
; GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
; CORRESPONDENCE LAW Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                          Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-280-805-145/c
; Sequence 145, Application US/09280805
; Patent No. 6184212,
; GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 100.0%; Preu. ...
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APPLICATION NUMBER: US/09/280,805
FILING BATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UNMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ISPH-0346
                                                                                                                                                         1084 ATTATAGCCTTAGTGAAGAA 1103
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                                                                                                                                                                                               20 ATTATAGCCTTAGTGAGAA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
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INFORMATION FOR SEQ ID NO: 144:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,29
REFERENCE/DOCKET NUMBER: 1
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7TP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-144
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                              US-09-280-805-144/c
                ANTI-SENSE:
US-09-280-805-143
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APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: BYPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
STREET: Mariton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 146, Application US/09280805
; Sequence 146, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2; TITLE OF INVENTION: EXPRESSION
; CORRESPONDENCE ADDRESS:
; ADDRESSER: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                          COUNTRY: NO
COUNTRY: NO
COUNTRY: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
COMPUTER: IBM PC
COMPUTER: IBM PC
COMPARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY, AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 15PH-0346
TELLCOMMUNICATION INPORMATION:
TELLECOMMUNICATION INPORMATION:
TELLECOMMUNICATION INPORMATION:
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STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IDSACIIE, 5.5 INCH, COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOCTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1100 AGAAGGACAAGAACTCTCAG 1119
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 145:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           U.S.A.
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ZIP: 08053
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 147, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 27
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
; CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: U.S.A.
ZIE: 08053
ZIE: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
COMPUTER: IN PC
COMPUTER: WINDOWS 95
SOFTWARE: WORDPERRECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NOMBER: U$/09/280,805
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match

0.8%; Score 20; UB
Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
PRIOR APPLICATION DATA:

PRIOR APPLICATION NUMBER: 09/048,810

FILING DATE: MAICH 26, 1998

ATTORNEY/AGENT INFORMATION:

NAME: Licata, Jane Massey

REGISTRATION NUMBER: 32,257

REFERENCE/DOCKET INFORMATION:

TELECOMMUNICATION INFORMATION:

TELEFANE: 609-810-1515

TELEFANE: 609-810-1454

INFORMATION FOR SEQ ID NO: 146:

SEQUENCE CHARACTERISTICS:

LENGTH: 20 base pairs

TYPE: Nucleic Acid

STRANDENESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1105 GACAAGAACTCTCAGATGAA 1124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TATOMAGE LICETE, JOHN MARREN JATOMAN MARKEL LICETE, JOHN MARREN JAZES REGISTRATION NUMBER: 1SPH-1FELECOMMUNICATION INFORMATION: TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1515
TERCHARACTERISTICS: LENGHALION FOR SEQ. ID NO: 147: SEQUENCE CHARACTERISTICS: LENGHALION FOR SEQ. Base pairs TYPE: Nucleic Acid STRANDEDNESS: Single TOPOLOGY: Linear TOPOLOGY: Linear HONG. 109-280-805-147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 GACAAGAACTCTCAGATGAA 1
                                                                                                                                                                                                                                                                                                                                                                           ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-280-805-147/c
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                                                                                                                                                                                                                                                                     Sequence 148, Application US/09280805
Factor No. 6184212
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION:
TITLE OF INVENTION:
CORRESPONDENCES. 271
CORRESPONDENCES. 271
CORRESPONDENCES. 271
CORRESPONDENCES. 120
STREET: 66 East Main Street
CORTY: Mailton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
     Length 20;
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US-09-280-805-149/c
i Sequence 199, Application US/09280805
j Patent No. 6184212
j GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2;
TITLE OF INVENTION: EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPTRY: NJ

ZIP: 08053

COMPUTER: NJ

COMPUTER READALE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1454
Query Match
0.8%; Score 20; DB 1; I
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                         1115 CTCAGATGAAGATGAGG 1134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1124 AGATGATGAGGTATATCAAG 1143
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 148: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs TYPE: Nucleic Acid STRANDEDNESS: Single TOPOLOGY: Linear ANTI-SENSE: Yes
                                                                                                                                                         20 crcagargargargagg 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 AGATGATGAGGTATATCAAG 1
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Gaps

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RESULT 304
US-09-280-805-151/C
; Sequence 151, Application US/09280805
; Patent No. 6164212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKT NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
NAME: Licata, Jane Massey
TELECOMMUNICATION INFORMATION:
NAME: Licata, Jane Massey
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 150:
SEQUENCE CHARACTER.FRICS:
LENGTH: 20 Dase pairs
TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1149 GTGTATCAGGCAGGAGAG 1168
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                         ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-150
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: Linear
TI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: U.S.A. ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-280-805-151
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; Sequence 150, Application US/09280805
; Patent No. 6184212
; Patent No. 6184212
; Patent No. 6184212
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: ANTISRNE MODULATION OF HUMAN MDM2
; TITLE OF INVENTION: EXPRESSION
; CORRESPONDENCE ADDRESS:
; CORRESPONDENCE ADDRESS:
; STREET: 66 East Main Street
; CITY: Mariton
; STRIE: NJ
; COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                               COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION: DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCA 26,1998
ATTORNEY/AGENT INFORMATION:
CLASSIFICATION TOWNER: 09/048,810
FILING DATE: MARCA 26,1998
ATTORNEY/AGENT INFORMATION:
CLASSIFICATION: TOWNER: TOWNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTY OR OS 3

ZIP: 08053

COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
MINIOWS 95
                          CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC
OPERATING SYSTEM:
OPERATING SYSTEM:
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GTAT 1154
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APPLICATION NUMBER: 09/048,810
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFRENCE/DOCKET NUMBER: 1SPH-
TELECOMMUNICATION INPORMATION:
TELEPHONE: 609-810-1515
INPORMATION FOR SEQ ID NO: 149:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TENTER NUCLEIC Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 rararcaagrracrererar 1
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     NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-149
                                                                                                                                                          U.S.A.
                                                                                     SITY: NO.
STATE: NJ.
COUNTRY: U.S.
10.9
                                                                                                       Marlton
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RESULT 307
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            Gaps
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100.0%; Pred. No. 3.3e+02;
cive 0; Mismatches 0; Indels
         0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: NJ
COUNTRY: U.S.A.

ZIP: 08052
ZIP: 08052
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: U$/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELEFAX: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 509-810-1515
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| Sequence 153, Application US/09280805
| Patent No. 6184212
| GENERAL INFORMATION:
| APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
| APPLICANT: Graham, Brett P. Monia
| TITLE OF INVENTION: EXPRESSION
| TITLE OF INVENTION: EXPRESSION
| COMPRESPONDENCE 271
| COMPRESSION COMPRESS: ADMINISTRY OF HOMAN MDM2 100 PRESSEE: Law Offices of Jane Massey Licata
                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Breet P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
      0; Mismatches
                                                         1161 GGGGAGAGTGATACAGATTC 1180
                                                                                                                                                                                                        RESULT 305
18-09-280-805-152/c
Sequence 152, Application US/09280805
Patent No. 6184212
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                                                                                                                   20 GGGAGAGTGATACAGATTC 1
20; Conservative
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Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-152
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Marlton
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Gaps
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| Saquence 154, Application US/09280805
| Patent No. 6184212
| GENERAL INFORMATION:
| APPLICANT: Graham, Brett P. Monia TITLE OF INVENTION: EXPRESSION ITLE OF INVENTION: EXPRESSION INVENTION: EXPRESSION INVENTION: EXPRESSION INVENTION: ADDRESSE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
| CITY: Marlton STATE: NJ COUNTRY: U.S.A.
| COUNTRY: U.S.A.
| COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
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3.3e+02;
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0.8%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                    SCHWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENI INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERNCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
INFORMATION FOR SEQ ID NO: 153:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: WORDPERPECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
                                                                                                                                                                                                                                                                                                                                                                                                          32,257
ER: ISPH-0346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1184 TGAAGAAGATCCTGAAATTT 1203
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 TGAAGAAGATCCTGAAATTT 1
                                                                                                                                                                      WINDOWS 95
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66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTI
COMPUTER: IBM PC
OPERATING SYSTEM: WII
                                                                                                                                                COMPUTER: IBM PC OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                            U.S.A.
                  Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-280-805-154/c
                                                          COUNTRY: U
ZIP: 08053
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STREET:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 155, Application US/09280805
Fatent No. 6184212
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTIESUSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CONTY: Marlton
                                                                                                                                                                                                                                                                                                               0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPPRATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.0%; Fred. No. 3.3
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION:
APPLICATION NUMBER: 09/048,810
ATTONNEY,AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECHONE: 609-810-1515
                         ISPH-0346
                                                                                                                                                                                                                                                                                                                                                                                                       1196 TGAAATTTCCTTAGCTGACT 1215
                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 reaaarrrccrragcreacr 1
REGISTRATION NUMBER: 32,257
                    REFERENCE/DOCKET NUMBER: IS
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: 009-810-1454
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                              TELEPAT. 609-810-1515
TELEPAT. 609-810-15454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
LENGTH: Nucleic Acid
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                       ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-154
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: Linear
TI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-280-805-155/c
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COUNTRY: U.
ZIP: 08053
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US-09-280-805-155
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Gaps
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     Sequence 156, Application US/09280805
; Patent No. 6184212
; GENERALI INFORMATION:
    APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
    APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
    APPLICANT: Graham, Brett P. Monia
    TITLE OF INVENTION: EXPRESSION
    INTRE OF INVENTION: EXPRESSION
    NUMBER OF SEQUENCES: 27;
    CORRESPONDENCE ADDRESS:
    ADDRESSEE: Law Offices of Jane Massey Licata
    STREET: 66 Bast Main Street
    CONTY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 20; DB 1; Length 20; 00.0%; Pred. No. 3.38+02;
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| Patent No. 6184212
| GENERAL INFORMATION:
| APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia | TITLE OP INVENTION: ANTISENSE MODULATION OF HUMAN MDM2 | TITLE OF INVENTION: EXPRESSION | NUMBER OF SEQUENCES: 271 | CORRESPONDENCE ADDRESS: Law Offices of Jame Massey Licata | STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPREFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 15PH-0346
TELECOMMUNICATION INFORMATION:
TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 156:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TTYPE: Nucleic Acid
STERBANDENCE CACID
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Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1220 GAAATGCACTTCATGCAATG 1239
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: Linear
HANTI-SENSE: Yes
US-09-280-805-156
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US-09-280-805-156/c
                                                                                                                                                                                                                                                                                                                                           STATE: NJ
COUNTRY: U.S
ZIP: 08053
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TELEFAX: 609-810-1454
; INFORMATION FOR SEQ ID NO: 1
; SEQUENCE CHARACTERISTICS:
; LENGTH: 20 base pairs
; TYPE: Nucleic Acid
; STRANNEDNESS: Single
; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-158
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-280-805-159/c
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       TELEPHONE:
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US-09-280-805-159
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Sequence 158, Application US/09280805

Patent No. 6184212

GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
NUMBER OF SEQUENCES: 271

CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ

COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.38+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                   ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IEM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 08053
CMONDTER READABLE FORM:
CMEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                        COMPUTER: IBM PC
CORPUTER: IBM PC
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIPICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26,1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jame Massey
REGISTRATION NUMBER: 12,257
REFERENCE/DOCKET NUMBER: 12,257
REFERENCE/DOCKET NUMBER: 15PH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: 609-810-1515
TELEFAX: 609-810-155:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 Dasse pairs
INFORMATION FOR SEQ ID NO: 157:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 Dasse pairs
TELEFAX: 100-810-1454
STRANDEDNESS: SINGle
THORNALICATION FOR SEQ ID NO: 157:
STRANDEDNESS: SINGle
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1226 CACTTCATGCAATGAATGA 1245
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APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 CACTTCATGCAATGAATGA 1
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W. U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TOPOLOGY: Linear; ANTI-SENSE: Yes US-09-280-805-157
COUNTRY: U.S.A.
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                                                                                                                                                                                                                                                                                    Sequence 159, Application US/09280805
Fatent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Grahm, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
INTEL OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
  Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR PAPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INPORMATION:
NAME: Licata, Jame Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFRONE 609-810-1454
Query Match
0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
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0.8%; Score 20; DB 1; Li
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                         1257 CCATCACATTGCAACAGATG 1276
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                                                                                                                                                            20 ccarcacarrecaacacare 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 159: SEQUENCE CHARACTERISTICS:
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1283 CCTTCGTGAGAATTGGCTTC 1302
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                                          WINDOWS 95
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TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: 162:
                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 609-810-1454
INFORMATION FOR SEO ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity 100.(
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      609-810-1454
                  COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         U.S.A.
MEDIUM TYPE:
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ZIP: 08053
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                                            ; Sequence 160, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
    APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
    APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
    APPLICANT: Graham, Brett P. Monia
    TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
    TITLE OF INVENTION: EXPRESSION
    NUMBER OF SEQUENCES: 271
    CORRESPONDENCE ADDRESS:
    ADDRESSEE: Law Offices of Jane Massey Licata
    STREET: 66 East Main Street
    CITY: Martton
    STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 161, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
   APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
   APPLICANT: Graham, Brett P. Monia
   TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
   TITLE OF INVENTION: RYPRESSION
   NUMBER OF SEQUENCES: 271
   CORRESPONDENCE ADDRESS:
   ADDRESSEE: Law Offices of Jane Massey Licata
   STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Licata, Jane Massey
REGISTRATION UNDRER: 32,257
REFRENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INPORMATION:
TELEPHONE: 609-810-1515
INPORMATION FOR SEQ ID NO: 160:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PECCOMPUTER: COMPUTER: IBM PECCOMPUTER: SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Herewith
CLASSIFFCATION:
PRIOR APPLICATION DATA:
RPICATICATION NUMBER: 09/048,810
FILING DATE: MATCH 26, 1998
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 08053
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-160
                                                                                                                                                                                                                                                                                                                                                           U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         U.S.A.
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STATE: NJ
                                     US-09-280-805-160/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-280-805-161/c
                                                                                                                                                                                                                                                                                                                                                         COUNTRY: U
ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:
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Gaps
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100.0%; Pred. No. 3.3e+02;
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Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 1SPH-0346
DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              100.0%; Pred. ...
                                                           SCHWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELERHONE: 609-810-1515
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APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
MUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata
STREET: ABAILON
STATE: NJ
Sequence 164, Application US/09280805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 AAAGGGAAAGATAAAGGGGA 1
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TELEFA: 609-810-1454
INFORMATION FOR SEQ ID NO: 164:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity 100.0
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: Linear ANTI-SENSE: Yes
                                         Patent No. 6184212
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U.S.A. ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          JS-09-280-805-165/c
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                                                                                                                                                                                                                                                                                          0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  18-09-280-805-163/c
| Sequence 163, Application US/09280805
| Patent No. 6184212.
| GENERAL INFORMATION:
| APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
| APPLICANT: Graham, Brete P. Monia
| TITLE OF INVENTION: EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IDEN PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION
PRIOR APPLICATION
PRIOR APPLICATION
PRIOR APPLICATION
PRIOR APPLICATION
FILING APPLICATION:
APPLICATION

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CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: Ge East Main Street
CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ISPH-0346
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NAME: Licata Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1544
INFORMATION FOR SEQ ID NO: 163:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDENESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20 GAATTGGCTTCCTGAAGATA 1
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                                 ; LENGTH: 20 base pairs
; TYPE: Nucleic Acid
; STRANDEDNESS: Single
; TOPOLGY: Linear
; AVII-SENNE: Yes
US-09-280-805-162
                                                                                                                                                                                                                                                                                       Query Match 0.8*
Best Local Similarity 100.0
Matches 20; Conservative
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Matches 20; Conservative
   SEQUENCE CHARACTERISTICS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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Fatent No. 6184212
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: 271
CORRESPONDENCES: 271
CORRESPONDENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Mailton
COMPUTER: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC OPERATING SYSTEM: WINDOWS 95 SOFTWARE: WORDPERFECT 6.0 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805 FILING DATE: herewith CLASHFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Le
3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 20; DB 100.0%; Pred. No. 3.3
                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1311 AAAGGGAAAGATAAAGGGGA 1330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC OPERATING SYSTEM:
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US-09-280-805-164/c

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Gaps
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                                                                                                       Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
COMPUTER: IBM PC
SOFTHARE: WINDOWS 95
SOFTHARE: WORDERFEET 6.0
CURRENT APPLICATION DATA:
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100.0%; Pred. No. 3.3e+02;
iive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT IRFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                               ; Sequence 167, Application US/09280805; Patent No. 6184212; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-280-805-168/c
; Sequence 168, Application US/09280805
; Patent No. 6184212
                                                                                                                                                                                             1333 TCTCTGAGAAAGCCAAACTG 1352
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INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
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Best Local Similarity 100.
Matches 20, Conservative
STRANDEDNESS: Single
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                      ; TOPOLOGY: Linear; ANTI-SENSE: Yes
US-09-280-805-166
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TOPOLOGY: Linear; ANTI-SENSE: Yes
US-09-280-805-167
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Marlton
                                                                                                                                                                                                                                                                                                                             US-09-280-805-167/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE:
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US-09-280-805-166/c
; Sequence 166, Application US/09280805
; Patent No. 6184212
; Patent No. 6184212
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: EXPRESSION
; TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STREEI: C. CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053

COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
CMMPUTER: IBM PC
CMMPUTER: MINDOWS 95
      CURRENT APPLICATION DATA:

PILING DATE: NO. 1972.

PILING DATE: No. 1972.

CLASSIFICATION NUMBER: US/09/280,805

FILING DATE: NO. 1998

APPLICATION NUMBER: 09/048,810

FILING DATE: MARCH 26, 1998

ATTORNEY/AGENT INFORMATION:

NAME: Licata, Jane Massey

REGISTRATION NUMBER: 32,257

REFERENCE/DOCKET NUMBER: 1SPH-0346

TELECOMMULICATION INFORMATION:

TELEPHONE: 609-810-1515

INFORMATION FOR SEQ ID NO: 165:

SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs

TYPE: NUCLeic Acid

STRANDENNESS: Single

TOPOLOGY: Linear

ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: ISM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIPICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTOCNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ISPH-0346
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INFORMATION FOR SEQ ID NO: 166:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
WORDPERFECT 6.0
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Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                      CLASSIFICATION

PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 09/048,810

FILING DATE: March 26, 1998

ATTORNEY/AGENT INFORMATION:

NAME: Licata, Jane Massey

REGISTRATION NUMBER: 32,257

REFERENCE/DOCKET NUMBER: 1SPH-0346

TELECOMMUNICATION INFORMATION:

TELEPHONE: 609-810-1515

TELEPAX: 609-810-1544

INFORMATION FOR SEQ ID NO: 169:

SEQUENCE CHARACTERISTICS:

LENGTH: 20 base pairs

TYPE: Nucleic Acid

STRANDENESS: Single
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COMPUTER: IBM PC
COMPUTER: IBM PC
COMPANIE: WORDPRIFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION
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REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPRAX: 609-810-1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1368 GCTGAAGAGGGCTTTGATGT 1387
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                       herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: Linear
ANTI-SENSE: Yes
US-09-280-805-169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Marlton STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-280-805-170/c
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                                                                                                                                                                                                                                                                                                                                                                        COUNTRY: U.S.A.

ZIP: 08053

COMPUTER RADALLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERRECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Herwith
CLASSIFICATION:
PRIOR APPLICATION BATCH
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jan Masses: 1SPH-0346
ATTORNEY/AGENT NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1454
INFORMATION FOR SEQ ID NO: 168:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDRESS: Single
TYPE: Nucleic Acid
STRANDEDRESS: Single
STRANDEDRESS: Single
ANTI-SENSE: Yes
US-09-280-805-168
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 169, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
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COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
....TRR: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERPECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1358 CTCAACACAAGCTGAAGAGG 1377
                                                                                                                                                                                                                                                                                                   CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-280-805-169/c
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0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                               Sequence 170, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A.
ZIF: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
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TITLE OF INVENTION:
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Matches
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                                           Query Match

0.8%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 172, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
                                                                                                                                                                                                                                                                                                Sequence 171, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street CITY: Marlton STATE: NJ STA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ISPH-0346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-034
TELECOMMUNICATION INFORMATION:
                                                                                                                                        1401 AAAACTATAGTGAATGATTC 1420
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1412 GAATGATTCCAGAGAGTCAT 1431
                                                                                                                                                                        20 AAACTATAGTGAATGATTC 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 GAATGATTCCAGAGAGTCAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHUNE: 009-810-1454
TELEPHUNE: 009-810-1454
INFORMATENA: 008-810-1454
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                              NUMBER OF SEQUENCES: 27
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-171
                                                                                                                                                                                                                                                       RESULT 324
US-09-280-805-171/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-280-805-172/c
US-09-280-805-170
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: MAILON.
STATE: MAILON.
ZID. 08053
ZOUNTRY: U.S.A.
ZID. 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
COMPUTER: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: DATE: WANDER: G.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: U$/09/280,805
FILING DATE: MARCh 26, 1998
ATTORNEY AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE, DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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COMDUTER READABLE FORM:
CMEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
              NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS: LAW Offices of Jane Massey Licata STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Apremith
CIASSIFICATION:
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IGAGG 1440
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EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1421 CAGAGAGTCATGTGT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Matches 20; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TOPOLOGY: Linear; ANTI-SENSE: Yes US-09-280-805-172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    U.S.A.
                                                                                               CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  S
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Gaps

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Indels

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Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0;
                                                                           1439 GGAAAATGATGATAAAATTA 1458
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                                                                                                                                                                                         RESULT 328
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                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.38+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S....
ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match

0.8%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORING ATTORING MARCH 26, 1998
ATTORING ATTORING MARCH 26, 1998
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1515
INFORMATION FOR SEQ ID NO: 173: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDENNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-280-805-174/c
; Sequence 174, Application US/09280805
; Patent No. 6184212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1434 GTTGAGGAAAATGATGATAA 1453
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 GTTGAGGAAATGATGATAA 1
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INFORMATION FOR SEQ ID NO: 174:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Linear
Yes
                                                                                                                                                                                                                                                                                                                          TOPOLOGY: Linear;
ANTI-SENSE: Yes
US-09-280-805-173
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; ANTI-SENSE:
US-09-280-805-174
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0.8%; Score 20; DB 1; Length 20;

Query Match

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Gaps
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                                                                                                                       Sequence 175, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-280-805-176/c

| Sequence 176, Application US/09280805
| Patent No. 6184212|
| Patent No. 6184212|
| GENERAL INFORMATION:
| APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Loren Brett P. Monia TITLE OF INVENTION: RAPISENSE MODULATION OF HUMAN MDM2 TITLE OF INVENTION: EXPRESSION NUMBER OF SEQUENCES: 271
| CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Licata, Jane Massey
RECISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1449 GATAAATTACACAAGCTTC 1468
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SEQUENCE GHARACTERISTICS.
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
ANTI-SENSE: Yes
20 GGAAATGATGATAAATTA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Marlton
                                                                                                       JS-09-280-805-175/c
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TOPOLOGY: Linear
ANTI-SENSE: Yes
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-280-805-178/c
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; Sequence 177, Application US/09280805
; Patent No. 6184312
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
; TITLE OF INVENTION: BYPRESSION
; NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
; STRIEE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                              ZIP: 08053
ZIP: 08053
ZIP: 08053
ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INPORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
FILEPHONE: 609-810-1515
FILEPHONE: 609-810-1515
INPORMATION FOR SEQ IO NO: TELEPHONE: G09-810-1515
INPORMATION FOR SEQ IO NO: SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STORE Nucleic Acid
STOROLGY: Linear
ANTI-SENSE: Yes
US-09-280-805-176
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION TOWER: 09/048,810
FILING DATE: MAXCh 26, 1998
ATTORNEY/AGENT INFORMATION:
ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
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                                                       STATE: NJ
COUNTRY: U.S.A.
7TP: 08053
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         STREET: 00 -
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US-09-280-805-177/c
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Fatent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPTRY:

COMPUTER:

COMPUTER:

MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER:

COMPUTE
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NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1515
INFORMATION FOR SEQ ID NO: 177:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
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REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1466 TTCACAATCACAAGAAGTG 1485
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INFORMATION FOR SEQ ID NO: 178:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
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TYPE: Nucleic Acid
STRANDEDNESS: Single
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REFISTRATION NUMBER: 32,257
REFISTRATION NUMBER: ISPH-0346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1499 GCCATCAACTTCTAGTAGCA 1518
                                                                                                                                                                                                                                                                                                                                                              NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-
TELECOMMUNICATION INFORMATION:
TELEFAX: 609-810-1515
INFORMATION FOR SEQ 1D NO: 180:
SEQUENCE CHARATERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
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Best Local Similarity 100.0
Matches 20; Conservative
                                                                    COMPUTER READABLE FORM:
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                          U.S.A.
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                                                                                            MEDIUM TYPE:
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US-09-280-805-180/c
; Sequence 180, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
; TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
; ADDRESSE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
                                                                                                                               US-09-280-805-179/c

Sequence 179, Application US/09280805

Patent No. 6184212

GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION 0.
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26,1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jame Massey
                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
1481 AAGTGAAGACTATTCTCAGC 1500
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                          20 AAGTGAAGACTATTCTCAGC 1
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INFORMATION FOR SEQ ID NO: 179:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.
Matches 20; Conservative
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; ANTI-SENSE: Yes
US-09-280-805-179
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COUNTRY: U.
ZIP: 08053
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| Patent No. 6184212
| GENERAL INFORMATION:
| APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
| APPLICANT: Graham, Brett P. Monia
| TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
| TITLE OF INVENTION: EXPRESSION
| NUMBER OF SEQUENCES: 271
| CORRESPONDENCE ADDRESS:
| ADDRESSEE: Law Offices of Jane Massey Licata
| STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
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COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM MINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
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              COMPUTER: IEM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURSENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFFCATION:
PRIOR PAPLICATION NUMBER: 09/048,810
FILING DATE: MAYON 26, 1998
ATTORNEY/AGENT INFORMATION:
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TELECOMMUNICATION INFORMATION:

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TOPOLOGY: Linear
TI-SENSE: Yes
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US-09-280-805-184/c
                      RESULT 336
US-09-280-805-183/c
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                                                                                                                                                                                                                                                                                                                         STATE:
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Pred. No. 3.3e+02;
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100.0%; Pred. No. 3.3e+02;
tive 0; Mismatches 0; IndelB
                                                                                                                                                                                                                                                                                 0; Indels
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Sequence 182, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
SOFTWARE: WORDPERRECT 6.0
CURRENT APPLICATION DATA:
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
                                                                                                                                                                                                                                  Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION

CLASSIFICATION DATA:

PRIOR APPLICATION DATA:

PRIOR APPLICATION DATA:

PRIORNEY/AGENT INPORMATION:

NAME: Licata, Jane Massey

REGISTRATION NUMBER: 32,257

REFERENCE/DOCKET NUMBER: 158PH-0346

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELEPAX: 609-810-1515

TELEPAX: 609-810-1634

INFORMATION FOR SEQ ID NO: 182:

LENGTH: 20 base pairs

TYPE: Nucleic Acid

STRANDENNESS: 51031e
                                                                                                                                                                                                                                                                                                                         1506 ACTICIAGIAGCATIATITA 1525
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1517 CATTATTATAGCAGCCAAG 1536
TELEPHONE: 609-810-1515
TELEPAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 181:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                    20 ACTICIAGIAGCATIATIA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 CATTATTATAGCAGCCAAG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-182
                                                                                                                                             ; TOPOLOGY: Linear; ANTI-SENSE: Yes
US-09-280-805-181
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 20; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE:
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Gaps

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US-09-280-805-186/c
; Sequence 186, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; TITLE OF INVENTION:
; TITLE OF INVENTION:
; TITLE OF INVENTION:
; TITLE OF INVENTION:
; OUNBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
; CITY: Marlton
; CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                             0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02;
                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                 Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                 1541 TGTGAAAGAGTTTGAAAGGG 1560
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1550 GITTGAAAGGGAAGAACCC 1569
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
; INFORMATION FOR SEQ ID NO: ;
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
; TYPE: Nucleic Acid
; TYPE: Nucleic Acid
; STRANDEDNESS: Single
; AMTI-SENSE: Yes
US-09-280-805-185
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 08053
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US-09-280-805-186
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY:
                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 340
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                   MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC
                                                                                                                                                                                                                                                                 ALICAMEAN JOINT NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 609-810-1515
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                   COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOCHWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFFCATION:
PRIOR APPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFRENCE/DOCKET NUMBER: ISPH-0346
FELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1533 CAAGAAGATGTGAAAGAGTT 1552
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 CAAGAAGATGTGAAAGAGTT 1
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  COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: Linear
HANTI-SENSE: Yes
US-09-280-805-184
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Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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Patent No. 6184212
GENERAL INFORMATION: APPLICANT: Graham, Brett P. Monia
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY:

ZIP: 08053

COMPUTER READDBLE PORM:
MEDIUM TYEE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE:
PRICK APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MATCH 26, 1998
ATTORNEY/AGENT INFORMATION:
                 CURRENT APPLICATION DATA:
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 1SPH-0346
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 609-810-1544
INFORMATION FOR SEQ ID NO: 188:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDENESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1566 ACCCAAGACAAAGAAGAGAG 1585
         OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 ACCCAAGACAAAGAAGAGAG 1
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REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: IS
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TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Marlton
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Sequence 188, Application US/09280805

Sequence 188, Application US/09280805

Fatent No. 6184212

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.

APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: EXPRESSION

NUMBER OF SEQUENCES:

NUMBER OF SEQUENCES:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street

STREET: NJ

COUNTRY: NJ

COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                           APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2 TITLE OF INVENTION: EXPRESSION NUMBER OF SQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 20; DB 1; Lengard, No. 3.3e+02;
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COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                                                                                                                                                                                                                                        ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ISPH-0346
                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFFCATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY AGENT INFORMATION:
NAME: Licata, Jane Massey
US-09-280-805-187/c
; Sequence 187, Application US/09280805
; Datent No. 6184212
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1560 GAAGAACCCAAGACAAAGA 1579
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1St
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 187:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-280-805-187
                                                                                                                                                                                                                                                                                         CITY: N
STATE:
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BENERAL INFORMATION:
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                                                                                                                                   0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                   Sequence 190, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: ANTISENSE CORRESSION
NUMBER OF SEQUENCES: 271
CORRESSONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DIEKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 20; DB 1; Le
Pred. No. 3.3e+02;
                                                                                                                              Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1515
INFORMATION FOR SEQ ID NO: 190:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
FILING DATE: herewith
CLASSIPICATION:
                                                                                                                                                                                                                       1580 AGAGAGTGTGGAATCTAGTT 1599
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 344
US-09-280-805-191/c
; Sequence 191, Application US/09280805
; Patent No. 6184212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1605 CTTAATGCCATTGAACCTTG 1624
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                               20 AGAGAGTGTGGAATCTAGTT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 criaargccarrgaaccrrg 1
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0.8%; Sc
Best Local Similarity 100.0%; P:
Matches 20; Conservative 0;
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
ANTL-SENSE: Yes
US-09-280-805-189
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                        US-09-280-805-190/c
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Fatent No. 6184212
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
INTRER OF SEQUENCES: 27;
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 20;
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2 TITLE OF INVENTION: EXPRESSION NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                      MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 20; DB 1; Lot 100.0%; Pred. No. 3.3e+02;
                                                                                                                                          ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.0%; Pred. No. ...
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                   SCHWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
PELLING DATE: herewith
CLASSIFICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION:
APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
ATYCNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REJECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 10454
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                                                                                                                                                                                                                                                                                                                                                                    WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                             U.S.A.
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US-09-280-805-192/c
                                                                                                                                                                                                                                             COUNTRY: U
ZIP: 08053
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Gaps

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                                                                           Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTIERNE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Mailton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKT NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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; Sequence 195, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Lozen J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
                                                                                                                                                                                                                                                                                                                              US-09-280-805-194/c
; Sequence 194, Application US/09280805
; Patent No. 6184212
                                                                                                                                                                              1648 AAAATGGTTGCATTGTCCAT 1667
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INFORMATION FOR EXO ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: NJ
COUNTRY: U.S.A.
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                          US-09-280-805-193
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| Sequence 193, Application US/09280805
| Patent No. 6184212
| GENERAL INFORMATION:
| APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia
| TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2 TITLE OF INVENTION: RAPRESSION
| NUMBER OF SEQUENCES: 271
| CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 Bast Main Street
CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFRENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPAK: 609-810-1515
INFORMATION FOR SEQ 1D NO: 193:
SEQUENCE CHARACTERISTICS:
LENGAL: 20 Asse pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
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APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                 RAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFRENCE/DOCKET NUMBER: 1SPH-
TELECOMMUNICATION INFORMATION:
TELEPAX: 609-810-1515
INFORMATION FOR SEQ ID NO: 192:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDENESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              20 GTGTGATTTGTCAAGGTCGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.8
Best Local Similarity 100.
Matches 20; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: Linear
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                          FILING DATE: h
CLASSIFICATION:
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US-09-280-805-192
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AESULT 350
US-09-280-805-197/c

i Sequence 197, Application US/09280805
i Patent No. 61494212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITT: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Indels
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MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEBORHONE: 600-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                  NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1545
INFORMATION FOR SEQ ID NO: 196: SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
LENGTH: 20 base pairs
TYPE: Nucleic Acid
TYPE: Nucleic Acid
TYPE: STRANDENNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1675 CAGGACATCTTATGGCCTGC 1694
        PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
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; INFORMATION FOR SEQ ID NO: 197:
; SEQUENCE CHARACTERISTICS:
; LENGTH: 20 base pairs
; TYPE: Nucleic Acid
; STRANDEDNESS: Single .
; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-197
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-196
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| Patent No. 6184212
| GENERAL INFORMATION:
| GENERAL INFORMATION:
| APPLICANT: Graham, Brett P. Monia | TITLE OF INVENTION: RANTSENSE MODULATION OF HUMAN MDM2 | TITLE OF INVENTION: RANTSENSE NO | NUMBER OF SEQUENCES: 271 | CORRESPONDENCE ADDRESS: | CORRESPONDENCE ADDRESS: | CORRESPONDENCE ADDRESS: | STREET: 66 East Main Street | CITY: Marlton | STREET: 66 East Main Street | COUNTRY: U.S. | C.S. | C.S.
                                                                                                                                                                                                         CITY: Marlton
STREET: 66 Eabt Follow
STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
COMPUTER: RADABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION:
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100.0%; Pred. No. 3.3e+02;
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2 TITLE OF INVENTION: EXPRESSION NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS: ADDRESS: ADDRESS: ADDRESSE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC.
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CAPPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
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Matches 20; Conservative
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Gaps

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COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                         ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
                                                                                                                                                                                                                      COMPUTER: IMP FC
COMPUTER: IMP FC
COURENT APPLICATION DATA:
APPLICATION NATA:
APPLICATION NATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
RAPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEBRONE: 000011151
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 20 base pairs
TYPE: Nucleic Acid
       CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TOPOLOGY: Linear; ANTI-SENSE: Yes US-09-280-805-199
                                                                                                                   U.S.A.
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                                                                      Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS:
                                                                                                                   COUNTRY: U
ZIP: 08053
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ZIP: 08053
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                                                                                                STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                        US-09-280-805-198/c
; Sequence 198, Application US/09280805
; Patent No. 6184312
; Patent No. 6184312
; Patent No. 6184312
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
; TITLE OF INVENTION: EXPRESSION
; CORRESPONDENCE ADDRESS:
; CARRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; CITY: Mailton
; CITY: Mailton
Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; Live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 199, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM FOR
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1454
INFORMATION FOR SEQ ID NO: 198:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1690 CCTGCTTTACATGTGCAAAG 1709
                                                                                           1684 TTATGGCCTGCTTTACATGT 1703
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
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Best Local Similarity 100.0
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-280-805-199/c
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US-09-280-805-198
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                                                                                                                        0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; Live 0; Mismatches 0; Indels
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MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPPRATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
                                                     1702 GTGCAAAGAAGCTAAAGAAA 1721
                                                                    20 GTGCAAAGAAGCTAAAGAAA 1
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1720 AAAGGAATAAGCCCTGCCCA 1739
                                                                     20 AAAGGAATAAGCCCTGCCCA 1
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; ANTI-SENSE: Yes
US-09-280-805-202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.C....

ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM:
OPERATING SYSTEM:
OPERATING SYSTEM: WANDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION 100
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAMME: Licate, Jane Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 132,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 132,257
REGISTRATION NUMBER: 150H-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Breter P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
                                                                     ISPH-0346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-280-805-201/c
; Sequence 201, Application US/09280805
; Patent No. 6184212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1710 AAGCTAAAGAAAAGGAATAA 1729
                 NAME: Licata, Jane Massey
REGISTRATION UNUBER: 32,257.
REFERENCE/DOCKET NUMBER: 15PH-
TELECOMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 200: SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
TOPOLOGY: Linear
US-09-280-805-200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 AAGCTAAAGAAAAGGAATAA 1
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INFORMATION FOR SEQ ID NO: 201:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.8
Best Local Similarity 100.
Matches 20; Conservative
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STATE: NJ
COUNTRY: U.
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US-09-280-805-201
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US-09-203-203/c
US-09-208-203. Application US/09280805
; Sequence 203. Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: LOTEN J. Miraglia, Pamela Nero, Mark J.
APPLICANT: LOTEN J. MIRSENSE MODULATION OF HUMAN MDM2; TITLE OF INVENTION: EXPRESSION
; TITLE OF INVENTION: EXPRESSION; NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 20;
'18-09-280-805-202/c
'Sequence 202, Application US/09280805
'Sequence 202, Application US/09280805
'Sequence 202, Application US/09280805
'GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
'TITLE OF INVENTION: EXPRESSION
'NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:

ZIP: 0853

COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: DIM PORTE: DISKETTE, 0.0

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Lideata, Jane Massey
REGISTRATION NUMBER: 32,257
                                                                                                                                                                                                 CITY: Mariton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REGISTRATION NUMBER: 32,257
REPERENCE/DOCKET UMBER: ISPH-0346
TELECOMMUNICATION: TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1726 ATAAGCCCTGCCCAGTATGT 1745
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INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
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1757 TCAAATGATTGTGCTAACTT 1776
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ZIP: 08053
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2 TITLE OF INVENTION: EXPRESSION NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: NG.
COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
COMPUTER: WORDERRECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
                                                           ZIP: 08053

ZIP: 08053

COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
                                                                                                                                                                                             SOFTWARE: WORDPREKET 6.0

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 12,257
REPERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFAX: 609-810-1515
TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 203:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 204, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; TOPOLOGY: Linear; ANTI-SENSE: Yes US-09-280-805-203
                                            U.S.A.
Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-280-805-204/c
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Sequence 205.4 Application US/09280805

Patent No. 6184212

GENERAL INFORMATION:

APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: EXPRESSION

NUMBER OF SEQUENCES: 27

CORRESPONDENCE ADDRESS:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street

CONTY: Marlton
                                                                                                                                                                                                                                                                                                                     0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IAM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jame Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 15PH-0346
TELECOMMUNICATION INFORMATION:
NAME: Licata, Jame Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 15PH-0346
TELECOMMUNICATION INFORMATION:
TELEBHONE: 609-810-1515
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPRONE: 609-810-1515
TELEPRAX: 609-810-1454
TELEFAX: 609-810-1464
INFORMATION FOR SEQ ID NO: 204:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TYPE: TOPOLOGY: Linear
STRANDENNES: Yes
US-09-280-805-204
                                                                                                                                                                                                                                                                                                                                                                                                                        1745 TAGACAACCAATTCAAATGA 1764
                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 TAGACCAATTCAAATGA 1
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INFORMATION FOR SEQ ID NO: 205:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                Ouery Match
Best Local Similarity 100.0
Matches 20, Conservative
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ANTI-SENSE: Yes
US-09-280-805-205
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; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
; VUMBER OF SEQUENCES: 27;
; CORRESPONDENCE ADDRESS:
; ADDRESSE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
; CITY: Marlton
                                                               GENERAL INPORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Breatt P. Monia
APPLICANT: Graham, Breatt P. Monia
APPLICANT: Graham, Breatt P. Monia
TITLE OF INVENTION: EXPRESSION
NUMBER OF EXQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STAFE: M.
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 20; DB 1; Length 20;
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                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A.

ZIP: 08053
COMPUTER READBALE FORM:
MEDIUM TYER: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/48,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OF 206:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPER: Nucleic Acid
STRANDEDMESS: Single
TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pred. No. 3.3e+02;
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0.8%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
US-09-280-805-206/c
; Sequence 206, Application US/09280805
; Patent No. 6184212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1787 GTTGACCTGTCTATAAGAGA 1806
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 GTTGACCTGTCTATAGAGA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ઠે
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COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Le
3.3e+02;
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0.8%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                       CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith

CLASSIFICATION DATA:

APPLICATION NUMBER: 09/048,810

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 09/048,810

FILING DATE: March 26,1998

ATTONNEY/AGENT INFORMATION:

NAME: Licata, Jane Massey

REGISTRATION NUMBER: 15PH-0346

TELECOMMUNICATION INFORMATION:

TELEPHONE: 609-810-1515

TELEPHONE: 609-810-155

TELEPHONE: CHARACTERISTICS:

LENGTH: 20 base pairs

TYPE: Nucleic Acid

TYPE: Macleic Acid
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APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
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OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1798 TATAAGAGAATTATATTT 1817
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 TATAAGAGAATTATATATT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-207
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STATE: NJ
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100.0%; Pred. No. 3.3e+02;
                                                                                                                                                                                           Score 20; DB 1; Length 20;
Pred. No. 3.3e+02;
                                                                                                                                                                                                                             0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                           Sequence 209, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREE:
CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Prec. ...
                                                                                                                                                                                     Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ISPH-0346
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: WORDFERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                                                                                                                                                                                                                1804 AGAATTATATATTTCTAACT 1823
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
PRIOR APPLICATION DATA:
PPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                     20 AGAATTATATATTTCTAACT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 TTATATATTTCTAACTATAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 15F
TELECOMMUNICATION:
TELEPHONE: 609-810-1515
                           208:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 009-810-1454
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TELEFAX: 609-810-1454
INPORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANEEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.4
Matches 20; Conservative
                                                                                                             TOPOLOGY: Linear
HAVII-SENSE: Yes
US-09-280-805-208
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                            US-09-280-805-209/c
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Gaps
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             Sequence 210, Application US/09280805

Patent No. 6184212

GENERAL INFORMATION:

APPLICANT: LOREN J. Miraglia, Pamela Nero, Mark J.

APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2

TITLE OF INVENTION: 27PL

CORRESPONDENCE ADDRESS:

ADDRESSES: Law Offices of Jane Massey Licata

STREET: 66 East Main Street

COLLY: Marlton
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; Sequence 211, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
    APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
    APPLICANT: Graham, Brett P. Monia
    TITLE OF INVENTION: EXPRESSION
    TITLE OF INVENTION: EXPRESSION
    NUMBER OF SEQUENCES: 271
    CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 08053
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
COMPUTER: IBM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMMUTER READBALE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1; Le
3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  32,257
32,257
70. ISPH-0346
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: IS
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 210:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     609-810-1454
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: U.S.A. ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STATE: NJ
COUNTRY: U.8
JS-09-280-805-210/c
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                 US-09-280-805-213/c
; Sequence 213. Application US/09280805
; Patent No. 6184212
; Patent No. 6184212
; General Information:
    APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
    APPLICANT: Graham, Brett P. Monia
    TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
    TITLE OF INVENTION: EXPRESSION
    NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
    ADDRESSEE: Law Offices of Jane Massey Licata
    STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 20;
                                                                                                                                                         Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1; Le
. 3.3e+02;
                                                                                                                                                         Query Match 0.8%; Score 20; DB 1; L. Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERPECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MATCH 26, 1998
ATTORNEY AGENT INFORMATION:
NAME: Licata, Jame Massey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPAX: 609-810-1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 367
US-09-280-805-214/c
; Sequence 214, Application US/09280805
                                                                                                                                                                                                                                            1832 CTAGGAATTTAGACAACCTG 1851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1840 TTAGACCAGCAGAATTTAT 1859
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 213:
SEQUENCE CHARACTREISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDMESS: Single
TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                      20 CTAGGAATTTAGACAACCTG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 Tragacaaccrgaaarrrar 1
    ; LENGTH: 20 base pairs
; TYPE: Nucleic Acid
; STRANDEDNESS: Single
; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U.S.A. ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-280-805-213
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: U.S.A.
ZIP: 08053.
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
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SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ISPH-0346
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                                                                                                                                                                                                                                                                                                                                           TELEFAX: 609-810-1454

INFORMATION FOR SEQ ID NO: 211:
SEQUENCE CHARACTREISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
JOPOLOGY: Linear
US-09-280-805-211
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      20 TATATAACCCTAGGAATTTA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET UMBER: 151
TELECOMMUNICATION
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Mariton
STATE: NJ
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US-09-280-805-212/c
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APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
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                                                                                                                                                                                                                                                                                                       TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
CURRENT APPLICATION DATA:
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Best Local Similarity 100.
Matches 20; Conservative
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TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: Linear ANTI-SENSE: Yes US-09-280-805-215
                                                                       CLASSIFICATION:
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; Sequence 215, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: LOTEN J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: EXPRESSION
; TITLE OF INVENTION: EXPRESSION
; CORRESPONDENCE ADDRESS:
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
; COUNTRY: NJ
; COUNTRY: NJ
              GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
STREET: MA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
tive 0; Mismatches 0; Indels
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MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
COPERATING SYSTEM: WINDOWS 95
SOPTWARE: WORDPERFECT 6.0
                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                    SOFTWARE: WORDERFECT 6.0

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELERAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 214:
SEQUENCE CHARACTER FSITCS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1850 TGAAATTTATTCACATATAT 1869
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Best Local Similarity 100.0
Matches 20; Conservative
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US-09-280-805-214
Patent No. 6184212
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 08053
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US-09-280-805-216/c
; Sequence 216, Application US/09280805
; Patent No. 6184212
; Patent No. 6184212
; PAPPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
; TITLE OF INVENTION: EXPRESSION
; CORRESPONDENCE ADDRESS:
; CARRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; CITY: Malton
; CITY: Malton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTEY: U.S.A.

ZIP: 08053
COMPUTEY: U.S.A.
COMPUTER: READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDFREECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 20,1998
ATTORNEY AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 15SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REBERRENCE/DOCKET NUMBER: ISPH-03*
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      . 1855 TTTATTCACATATATCAAAG 1874
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NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REPRENCE/DOCKET NUMBER: ISPH
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             i LENGTH: 20 base pairs
; TYPE: Nucleic Acid
; STRANDEDNESS: Single
; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-218
APPLICANT: Graham, Brett
TITLE OF INVENTION: ANTI
TITLE OF INVENTION: EXPR
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: 609-810-1454
INFORMATION FOR SEQ 1D NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                         U.S.A.
                                                                                                                                                          CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   JS-09-280-805-219/c
                                                                                                                                                                                                         COUNTRY: U ZIP: 08053
                                                                                                                  ADDRESSEE:
                                                                                                                                         STREET:
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                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: LOVEN J. Miraglia, Pamela Nero, Mark J.
APPLICANT: LOVEN J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
INTRE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 27;
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: 0. COUNTRY: U.S.A.
                                                                                           Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY, AGENT INFORMATION:
NAME: Licata, Jame Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 15PH-0346
TELECOMMUNICATION NUMBER: 15.757
REFERENCE/DOCKET NUMBER: 15.757
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                                                                                        Score 20; DB 1; Let Pred. No. 3.3e+02;
                                                                                    Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                             US-09-280-805-217/c
; Sequence 217, Application US/09280805
; Patent No. 6184212
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US-09-280-805-218/c
; Sequence 218, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
                                                                                                                                                                            1865 TATATCAAAGTGAGAAATG 1884
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1872 AAGTGAGAAAATGCCTCAAT 1891
                                                                                                                                                                                                                         20 TATATCAAAGTGAGAAATG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     20 AAGTGAGAAATGCCTCAAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 217:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 20 base pairs
TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS: SIL

TOPOLOGY: Linear

ANTI-SENSE: Yes

US-09-280-805-217
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Matches 20; Conserve
                ; ANTI-SENSE:
US-09-280-805-216
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Sequence 219, Application US/09280805
Fatent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
ITILE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEB: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STREET: 66 East Main Street
COMPUTRY: U.S.A.
ZIP: 08053
COMPUTER: ISP PC
SECURATION: WINDOWS 95
SCOMPUTER: ISP PC
SECURATION: APPLICATION DATE:
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
Brett P. Monia
ANTISENSE MODULATION OF HUMAN MDM2
EXPRESSION
                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3.3e+02;
                                                                                                                        iB: Law Offices of Jane Massey Licata
66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.8%; Score 20; DB 1;
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Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ISPH-0346
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; TOPOLOGY: Linear; ANTI-SENSE: Yes US-09-280-805-221
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US-99-280-805-220/C
is 5equence 220, Application US/09280805
is Fatent No. 6184212
is GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
ITILE OF INVENTION: ARTISENSE MODULATION OF HUMAN MDM2
ITILE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCES: 271
CORRESPONDENCES: 271
STREET: Law Offices of Jane Massey Licata
STREET: Amailton
STREET: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
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MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION NUMBER: 09/048,810
FILING DATE: MARCH 26,1998
ATTORNEY/AGRAT INFORMATION:
NAME: Licata, Jane Massey
REFERENCY/DOCKET NUMBER: 12,157
REFERENCY/DOCKET NUMBER: 12,157
REFERENCY/DOCKET NUMBER: 12PH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECHONR: 609-810-1515
                                                                                                                                                                                                                                                                                                        ISPH-0346
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REPERENCE/DOCKET NUMBER: 1SPH-03
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1454
; SEQUENCE CHARACTERISTICS:
***CHARACTERISTICS:
***TELEPHONE: CHARACTERISTICS:
****TELEPHONE: CHARACTERISTICS:
****TELEPH
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; INFORMATION FOR SEQ ID NO: 220:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-220
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: Linear
ANTI-SENSE: Yes
US-09-280-805-219
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; Sequence 221, Application US/09280805
; Patent No. 6184212
; Patent No. 6184212
; Patent No. 6184212
; Patent No. 6184212
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
; TITLE OF INVENTION: EXPRESSION
; CORRESPONDENCE ADDRESS:
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; CITY: Marlton
; CITY: Marlton
Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.38+02;
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% Sequence 222, Application US/09280805

% Patent No. 184212.

GENERAL INFORMATION:

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.

APPLICANT: Graham, Breft P. Monia

TITLE OF INVENTION: EXPRESSION

TITLE OF INVENTION: EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 08053

ZIP: 08053

COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
COMPUTER: WINDOWS 95
SOFTWARE: WORDERFECT 6.0

CURRENT APPLICATION DATA:
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match

Query Match

Best Local Similarity 100.0%; Pred. No. 3.3

Matches 20; Conservative 0; Mismatches
                                                                                                       1898 AGATTTCTTCTTTAGTAT 1917
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26,1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                            20 AGATTTCTTCTCTTTAGTAT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: IS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: Nucleic Acid
STRANDEDNESS: Single
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Fatent No. 6184312
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Mailton
                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCAET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELESPENSE: 609-810-1815
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FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 12,257
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1454
INFORMATION FOR SEQ ID NO: 223:
SEQUENCE GHARACTERISTICS:
LENGER GHARACTERISTICS:
LENGER: Nucleic Acid
STRANBDENES: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 AGTATAATTGACCTACTTTG 1
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Matches 20; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                              ; TOPOLOGY: Linear; ANTI-SENSE: Yes
US-09-280-805-223
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US-09-280-805-224/c
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; Sequence 223, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loraham, Brett P. Monia
; TITLE OF INVENTION: RATISENSE MODULATION OF HUMAN MDM2; TITLE OF INVENTION: EXPRESSION
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
                                                                                                                                   STFALE.

COUNTRY: U.S.A.

ZIP. 08053

COMPUTER READABLE FORM:

MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE

NEDIUM TYPE: IN PC

WINDOWS 95
  NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                    COMPUTER: ILEN FC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION: PRICH 200,005
FILING DATE: herewith
CLASSIFICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26,1998
ATTORNEY/AGENT: NPORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELEPAN: 609-810-1155
TELEPAN: 609-810-1155
TELEPAN: 20 DASS DINO: 222:
SEQUENCE CHARACTERISTICS:
LINFORMATION FOR SEQ ID NO: 222:
SEGUENCE CHARACTERISTICS:
LENGTH: 20 DASS pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1908 TCTTTAGTATAATTGACCTA 1927
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; ANTI-SENSE: Yes
US-09-280-805-222
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STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-280-805-223/c
                                                                                                                             STATE: NJ
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  0; Indels
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Sequence 226, Application US/09280805
Fatent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
                                                                                                                                                                 US-09-280-805-225/c

Sequence 225, Application US/09280805

Parent No. 6184212

GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 20; DB 1; Le
Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
  0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Licata, Jane Massey
REGISTRATION UNUBER: 32,257
REFRENCE/DOCKET NUBBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEFHONE: 609-810-1515
TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 225:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IEM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION UNDER: 09/048,810
APPLICATION NUMBER: 09/048,810
FILING DATE: MATCH 26,1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1933 GTAGTGGAATAGTGAATACT 1952
                                            1920 TTGACCTACTTTGGTAGTGG 1939
                                                                  20 TTGACCTACTTTGGTAGTGG 1
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  Conservative
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Yes
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                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; ANTI-SENSE:
US-09-280-805-225
20;
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  Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 227 227.
Sequence 227
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COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
COMPUTER: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
REPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER:
PILING DATE: March 26, 1998
ATTORNEY AGENT INPORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1940 AATAGTGAATACTTATA 1959
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            20 AATAGTGAATACTTACTATA 1
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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTER.STICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                                                  U.S.A.
Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-280-805-227/c
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                                                2
                                                                                       COUNTRY:
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REGISTRATION NUMBER: 32,257

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20 TATAATTTGACTTGAATATG 1
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US-09-280-805-229
                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
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; Sequence 228, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: LOYEN J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: EXPRESSION
; TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
; STREET: Mariton
; STREET: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match

0.8%; Score 20; DB 1; Length 20;
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                        0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: ENA PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFCATION:
RAPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY, GARNI INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELLCOMMUNICATION INFORMATION:
                      ISPH-0346
                                                                                                                                                                                                                                                                                                                                                                                            1948 ATACTTACTATATTTGACT 1967
           REFERENCE/DOCKET NUMBER: ISPH.
TELECOMOUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 227:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDENESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                     20 ATACTTACTATAATTTGACT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTER.ESTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                    Query Match 0.89
Best Local Similarity 100.(
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Linear
Yes
                                                                                                                                                                                                              TOPOLOGY: Linear

ANTI-SENSE: Yes

US-09-280-805-227
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ZIP: 08053
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Gaps
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100.0%; Pred. No. 3.3e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 230, Application US/09280805
Fatent No. 6184212
GENERAL INFORMATION:
GENERAL INFORMATION:
TITUE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
STREET: 66 East Main Street
                                        Sequence 229, Application US/09280805
Patent No. 6184312
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
TITLE OF INVENTION: EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
FILING DATE: herewith
                                                                                                                                                                                                     NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
PPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION:
TELECHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1969 GAATATGTAGCTCATCCTTT 1988
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INFORMATION FOR SEQ ID NO:

SEQUENCE CHARACTERISTICS:

'RNGTH: 20 base pairs
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Best Local Similarity 100.0
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: Linear
TI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                   U.S.A.
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LITY: Marlton
STATE: NT
RESULT 382
US-09-280-805-229/c
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1956 TATAATTTGACTTGAATATG 1975

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NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISI
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
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                  TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
    609-810-1515
                                                                                                                                                                                                                                  Ouery Match
Best Local Similarity 100.C
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.8
Best Local Similarity 100.
Matches 20, Conservative
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TYPE: Nucleic Acid
                                                                                                                                                    TOPOLOGY: Linear
HANTI-SENSE: Yes
US-09-280-805-231
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; ANTI-SENSE: Yes
US-09-280-805-232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Marlton
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    TELEPHONE:
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; Sequence 231, Application US/09280805
; Patent No. 6194212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
; TITLE OF INVENTION: EXPRESSION
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
; CITY: Marlton
; COUNTRY: U.S.
; COUNTRY: U.S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Score 20; DB 1; Lengt....
---d. No. 3.30+02;
0; Indels
                                    COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 15PH-0346
TELECOMMUNICATION INFORMATION:
TELEFAX: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 609-810-1515
TELEFAX: 609-810-164
INFORMATION FOR SEQ ID NO: 230: SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NDAER:
APPLICATION NDAER:
FILING DATE: MACCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licate, Jane Massey,
REGISTRATION NUMBER: 32,25,
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
                                                                                                    OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION UNBER: US/09/280,805
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILLING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1973 ATGTAGCTCATCCTTTACAC 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 Argradercarecriracae 1
                                                                                    COMPUTER: IBM PC OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-230
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC
OPERATING SYSTEM:
U.S.A.
                                                                                                                                                                                                                    CLASSIFICATION:
COUNTRY:
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| INFORMATION FOR EACH SEED IN No. 211;
| SEQUENCE CHARACTERISTICS: | InfoRMATION FOR EACH SEED IN No. 211;
| CHARACTERISTICS: | InfoRMATION: | InfoRMATION:
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; Sequence 235, Application US/09280805; Patent No. 6184212; GENERAL INFORMATION: GENERAL INFORMATION: APPLICANT: Graham, Brett P. Monia TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2: TITLE OF INVENTION: EXPESSION NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS: ADDRESSE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
; CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053.
ZIP: 08053.
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FTI:NG DATE: herewith
DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                          CURRENT APPLICATION DATA:

POSTERIN APPLICATION DATA:

PRILING DATE: Herewith

CLASSIFICATION:

PRIOR APPLICATION:

PRIOR APPLICATION:

PRIOR APPLICATION:

APPLICATION:

PRIOR APPLICATION:

APPLICATION NUMBER: 09/048,810

FILING DATE: March 26, 1998

ATTONNEY, AGENT INFORMATION:

NAWE: Lidata, Jane Massey

REGISTRATION NUMBER: 32,257

REFERENCE/DOCKT NUMBER: 169H-0346

TELEPHONE: 609-810-1515

TELEPHONE: 609-810-1515

INFORMATION FOR SEQ ID NO: 234:

SEQUENCE CHARACTERISTICS:

LENGTH: 20 base pairs

TYPE: Nucleic Acid

STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ISPH-0346
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 rrraaaraarrrcracrcrg 1
                                          WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: IST
TELECOMMUNICATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
                    COMPUTER: IBM PC OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 388
US-09-280-805-235/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-280-805-234
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Sequence 234, Application US/09280805

Sequence 234, Application US/09280805

Patent No. 6184212

GENERAL INFORMATION:

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia

TILLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2

TITLE OF INVENTION: EXPRESSION

NUMBER OF SEQUENCES: 271

CORRESPONDENCE ADDRESS:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 20;
                                                                                                                              APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: 271
CORRESPONDENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                    COMPUTRY: U.S.A.
ZIP: 08053
COMPUTRY: U.S.A.
COMPUTRY: BADABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIPICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/046,810
FILING DATE: MACCh 26, 1998
ATTORNEY/AGENT MACCh 26, 1998
ATTORNEY/AGENT MACCh 26, 1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                            Sequence 233, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1997 TCCTAATTTTAAATTTTC 2016
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEFAX: 609-810-1454
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 TCCTAATTTTAAATAATTC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ANTI-SENSE: Yes
US-09-280-805-233
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            U.S.A.
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STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
                                          US-09-280-805-233/c
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2051 TTTTCTTAAATATGTATATG 2070
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
Matches 20; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                         U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U
ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-280-805-237
                                                                                                                                                                                                                                                                         COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
                                                                                                                                                                    Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                                                                                                                                                                    Score 20; DB 1; Losted. No. 3.3e+02;
                                                                                                                                                              Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER:
APPLICATION NUMBER:
PLING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licate, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECHONE: 609-810-1515
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                    US-09-280-805-236/c
; Sequence 236, Application US/09280805
; Datent No. 618421;
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2020 TCTGTCTTAAATGAGAAGTA 2039
                                                                                                                                                                                                                                                2015 TCTACTCTGTCTTAAATGAG 2034
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 236: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs TYPE: Nucleic Acid STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                      20 rcracrcrcrcrraaargag 1
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; SEQUENCE CHARACTERISTICS:
; LENGTH: 20 base pairs
; TYPE: Nucleic Acid
; TYPE: NCOPLOGY: Linear
; AVTI-SENSE: Yes
US-09-280-805-235
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: Linear; ANTI-SENSE: Yes
US-09-280-805-236
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Gaps
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Sequence 237, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 391
US-09-280-805-238/c
; Sequence 238, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; TITLE OF INVENTION: Brett P. Monia
; TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 271
; CORRESPONDENCE Law Offices of Jane Massey Licata
; STREET: Law Offices of Jane Massey Licata
; STREET: Law Offices of Jane Massey Licata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                               ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MATCH 26, 1998
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAWE: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC
OPERATING SYSTEM:
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WINDOWS 95

US-09-280-805-237/c

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STRANDEDNESS: Single
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MEDIUM TYPE: DISKETT
                            ; TOPOLOGY: Linear; ANTI-SENSE: Yes US-09-280-805-239
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 239, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFFCATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAMM: Licata, Jane Massey
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION NUMBER: 1SPH-0346
TELECOMMUNICATION NUMBER: 1SPH-0346
TELECOMMUNICATION NUMBER: 1SPH-0346
TELEPRAN: 609-810-1515
TELEFRAX: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street CITY: Marlton STATE: NJ COUNTRY: U.S.A.
  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFTCATION:
CLASSIFTCATION:
PROOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY, AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION NUMBER: ISPH-0346
TELECOMMUNICATION NUMBER: ISPH-0346
TELECOMMUNICATION NUMBER: ISPH-0346
TELECOMMUNICATION NUMBER: 15PH-0346
TELECOMMUNICATION NUMBER: ISPH-0346
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLLARY
THERETOM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2059 AATATGTATATGACATTTAA 2078
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WORDPERFECT 6.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: Linear
ANTI-SENSE: Yes
US-09-280-805-238
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Sequence 240, Application US/09280805

Sequence 240, Application US/09280805

Patent No. 6184212

GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Grahm, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCES. 271
CORRESPONDENCES. 271
STREET: 66 East Main Street
CITY: Marlton
STATE: N.
COUNTRY: U.S.A.

COUNTRY: U.S.A.
  Length 20;
                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.8%; Score 20; DB 1; Length 20;
.00:0%; Pred. No. 3.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IEM PC OPERATING SYSTEM: WINDOWS 95
Score 20; DB 1; L
Pred. No. 3.3e+02;
Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Best Local Similarity 100:0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OPERATING SYSTEM: MINLOWS SO SOFTWARE: WARDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Herewith
CLASSIFICATION NUMBER: 09/048,810
FILING DATE: Herewith
FILING DATE: MARCH 26,1998
ATTORNEY/AGENT INFORMATION:
MAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-154
INFORMATION FOR SEQ ID NO: 240:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDENESS: Single
ANTI-SENSE: Yes
US-09-280-805-240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 394
US-09-280-805-241/c
; Sequence 241, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
                                                                                         2072 CATITAAAIGIAACTIAITA 2091
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2103 ACCGAGTCTTGCTCTGTTAC 2122
                                                                                                                                        20 CATTTAAATGTAACTTATTA 1
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US-09-280-805-243/c
; Sequence 243, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: EXPRESSION
; TITLE OF INVENTION: EXPRESSION
; UNMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESSS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; CITY: Marlton
; CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CCUNTRY: NO
CCUNTRY: NO
CCUNTRY: NO
ZIP: 08053
CMPUTER READABLE FORM:
MEDITUR TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: WINDOWS 95
SOFTWARE: WORDERREFOR 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Merewith
CLASSIFICATION NUMBER: US/09/280,805
FILING DATE: Merewith
TLASSIFICATION NUMBER: 109/048,810
FILING DATE: MARSEY
FILING DATE: MARSEY
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 1259
TRIEFRANGOCKET NUMBER: 158H-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-11515
TELEFAX: 609-810-1154
TELEFAX: 20 Dase pairs
TYPE: Nucleic Acid
STRANDENDESS: Single
TYPE: VENSES: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   100.0%; Prea. ...
                       CLASSIFICATION
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26,1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 15PH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPAX: 609-810-1545
INFORMATION FOR SEQ ID NO: 242:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2116 CTGTTACCCAGGCTGGAGTG 2135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Best Local Similarity 100. Matches 20; Conservative
       herewith
                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: Linear
ANTI-SENSE: Yes
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Fatent No. 618412
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Mailton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2 TITLE OF INVENTION: EXPRESSION
                                                                                                                                                                                                                                           ZIP: 08053

COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
CLASSIFICATION:
PRIOR APPLICATION:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 08053
CONFUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
                                                                                                                                    ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.: Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1515
INFORMATION FOR SEQ ID NO: 241:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/09/280,805
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NUMBER OF SEQUENCES: 2:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Office
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: Linear
TI-SENSE: Yes
                                                                                                                                                                                                                             U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-280-805-242/c
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US-09-280-805-241
                                                                                                                                                                                                      STATE: N. COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: N
STATE:
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PRIOR APPLICATION DATA:
    TITLE OF INVENTION:
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                                                                                                   Gaps
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                                               0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                        RESULT 397
US-09-280-805-244/c
Sequence 244, Application US/09280805
Fatent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2;
TITLE OF INVENTION: EXPRESSION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 245, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANI: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRA.
ZIP: 08053
COMPUTER: READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
CAMPUTER: IBM PC
TANDUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION D:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFRENCE/DOCKET NUMBER: ISPH-0346
TELEPHONE: 609-810-1515
                                                                                                                                           2123 CCAGGCTGGAGTGCAGTGGG 2142
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                                          Query Match
Best Local Similarity 100.(
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: Linear ANTI-SENSE: Yes US-09-280-805-244
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US-09-280-805-243
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Sequence 246, Application US/09280805

Patent No. 6184212

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION:
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                   NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STARE: NJ
COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                      COMPUTER: 15M PC.

OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WONDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY, AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REJERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2140 GGGTGATCTTGGCTCACTGC 2159
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EXPRESSION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                          COMPUTER: IBM PC OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-245
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OPERATING SYSTEM:
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Gaps

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Indels

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Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                RESULT 401
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Sequence 247, Application US/09280805

Patent No. 6184212

GENERAL INFORMATION:

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.

APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2

TITLE OF INVENTION: EXPRESSION

NUMBER OF SEQUENCES: 271

CORRESPONDENCE ADDRESS:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: DISCRIT:
MEDIUM TYPE: DISCRIT:
MEDIUM TYPE: DISCRIT:
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION OFFEWAITH
PRIOR APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licate, Jame Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
               FILING DATE: MARCH 26, 1998
ATTORNEY/AGENT INFORMATION:
MAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
FELECOMMUNICATION INFORMATION:
FELEPAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 246:
SEQUENCE CHARACTER.ESTICS:
LENGTH: 20 base pairs
FYPE: Nucleic Acid
STRANDEDNESS: Single
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
US-09-280-805-246
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2146 TCTTGGCTCACTGCAAGCTC 2165
  09/048,810
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INFORMATION FOR SEQ ID NO: 247:
SEQUENCE GHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
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MEDIUM TYPE: DISKETT
  APPLICATION NUMBER:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
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Length 20;

DB 1;

0.8%; Score 20;

Query Match

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Sequence 249, Application US/09280805

Patenty, no. 6184212

GENERAL INFORMATION:

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.

APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2;

TITLE OF INVENTION: EXPRESSION

NUMBER OF SEQUENCES: 271
                                                                                                                                                Sequence 248, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:

ZUP: 08053

COMPUTER READABLE FORM:
MBDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
TING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 20; DB 1; Le
100.0%; Pred. No. 3.3e+02;
tive 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
2153 TCACTGCAAGCTCTGCCCTC 2172
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      rccrg 2195
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
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                                         20 reacrecasecrerecere
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,29
REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2176 GGGTTCGCACCATTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         609-810-1454
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Best Local Similarity
Matches 20; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; TOPOLOGY: Linear; ANTI-SENSE: Yes US-09-280-805-248
                                                                                                                                                                                                                                                                                                                                                                                                Marlton
                                                                                                                                  US-09-280-805-248/c
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ADDRESSEE:

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Gaps
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US-09-280-805-251/c

Sequence 251, Application US/09280805

Patent No. 618412

GENERAL INFORMATION:

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.

APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2;

TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2;

COMPRESPONDENCE ADDRESS:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street

CITY: Marlton

CITY: Marlton

STATE: No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
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MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                       DB 1; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                Query Match

0.8%; Score zu; De Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
NAME: Licata, Jane Massey

REGISTRATION NUMBER: 32,257

REFERENCE/DOCKET NUMBER: 1SPH-0346

TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515

TELEPHONE: 609-810-1515

TELEPHONE: 609-810-1515

INFORMATION FOR SEQ ID NO: 250:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
LENGTH: 20 base pairs

LENGTH: 20 base pairs

TYPE: Nucleic Acid
STRANDEDNESS: Single

TOPOLOGY: Linear

NATI-SENSE: Yes
US-09-280-805-250
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDION TIPE: DISKETIE, 3.5 INCH, 1.44
COMPUTER: 1BM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION.:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTONNEY, AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: U.S.A. ZIP: 08053
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 250, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: Coren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESSONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                        ZIP: 0805 0.5.4.
ZIP: 0805 0.5.4.
ZIP: 0805 0.5.4.
ZIP: 0805 0.5.4.
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC
SOFTWARE: WINDDWS 95
SOFTWARE: WINDDWS 95
SOFTWARE: WINDDRESCT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: March 26, 1998
ATTORNEY/AGRY INFORMATION:
FILING DATE: March 26, 1998
ATTORNEY/AGRY INFORMATION:
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 32,257
RELECOMMUNICATION INFORMATION:
TELEPRAN: 609-810-1515
TELERAM: 609-810-1515
TELERAM: 609-810-1454
INFORMATION FOR SEQ ID NO: 249:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
  1E: Law Offices of Jane Massey Licata
66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2185 CCATTCTCCTGCCTCAGCCT 2204
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CLASSIFICATION:
PRIOR APPLICATION DATA: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
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Best Local Similarity 100.0
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: Linear

ANTI-SENSE: Yes

US-09-280-805-249
                                                                                             U.S.A.
           STREET: 00.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-280-805-250/c
                                                                     STATE: N. COUNTRY:
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COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION OF 810-1454
INFORMATION FOR SEQ ID NO: 253:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDENES: SINGLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION NATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
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REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
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HANTI-SENSE: Yes
US-09-280-805-253
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COUNTRY: U.S.A.
ZIP: 08053
                            U.S.A.
                            COUNTRY: U
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                                                                                                                                          Sequence 252, Application US/09280805
Fatent No. 6184212
GENERAL INPORMATION:
GENERAL INPORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CONTY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence 253, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
APPLICANT: Loranan, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
; CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CONTINENT TO THE TO CONTINENT WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MATCH 26, 1998
ATTORNEY/AGENT INFORMATION:
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2198 TCAGCCTCCCAATTAGCTTG 2217
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REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET UNBER: 1St
TELECOMMUNICATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.8
Best Local Similarity 100.
Matches 20; Conservative
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TI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                       U.S.A.
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US-09-280-805-252
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Gaps ö US-09-280-805-254/C
; Sequence 254, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: EXPRESSION
; TITLE OF INVENTION: EXPRESSION
; NUMBER OF SEQUENCES: 271
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; CITY: Marlton Length 20; COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95 COMPUTER READABLE FORM: MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE Query Match

0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;

TELECOMMUNICATION INFORMATION:

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COUNTRY: U.S.A. ZIP: 08053
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STATE: NJ
COUNTRY: U.S.A.
                                                   IS-09-280-805-256/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 255, Application US/09280805
Patent No. 6184212
GENERAL INFORMATION:
APPLICANT: LORD Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCE: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                         2213 GCTTGGCCTACAGTCATCTG 2232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2218 GCCTACAGTCATCTGCCACC 2237
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 GCCTACAGTCATCTGCCACC 1
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REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISI
TELECOMMUNICATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 255:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TELEPHONE: 609-810-1515
TELEPAX: 609-810-1515
INPORMATION FOR SEQ ID NO: 25.
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
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ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 20; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-280-805-255/c
                                                                                                                                                                 ; ANTI-SENSE:
US-09-280-805-254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-280-805-255
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COUNTRY:
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SEGULATION CONSTRUCTION US/92280805

SEGULATION COSTS, APPLICATION WITERIAL Pamela Nero, Mark J. APPLICANT: Locren J. Wiragilia, Pamela Nero, Mark J. APPLICANT: Carban J. Warragilia, Pamela Nero, Mark J. APPLICANT: Graham Breatt P. Monia TITLE OF INVESTION: EXPERSION MARS MATISHES MODULATION OF HUMAN MOME TITLE OF INVESTION: EXPERSION MARS MATISHES MODULATION OF HUMAN MOME TITLE OF INVESTION: EXPERSION MARS MATISHES MODULATION OF HUMAN MOME TITLE OF INVESTION: EXPERSION MARS MATISHES MODULATION OF HUMAN MOME MATISHES MODULATED ADDRESSED AND MATISHES MODULATION MARS. 158H-0346

STATE: NO CONTINUE MATISHES MODULATION MASS MODULATION MARS MODULATION MASS MODULATED ADDRESSED AND MATISHES MODULATION MASS MODULATED AND MASS MODULATION MASS MODULAT
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Gaps

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Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPERTY.

COMPUTER: NO

COMPUTER: READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGRY INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEBENOME: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.8%; Score 20; DB 1; L
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-280-805-259/c
; Sequence 259, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                             ; LENGTH: 20 base pairs
; TYPE: Nucleic Acid
; STRANDEDNESS: Single
; TOPOLGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-258
                                                   INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
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US-09-280-805-259
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US-09-200-0805-258/c
| Sequence 258, Application US/09280805
| Patent No. 6184212
| GENERAL INFORMATION:
| APPLICANT: Graham, Brett P. Monia
| TITLE OF INVENTION: RAPRESSION
| NUMBER OF SEQUENCES: 271
| CORRESPONDENCES: 271
| CORRESPONDENCES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
COMPUTER READABLE FORM:

MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: Herwith
CLASSIPICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licate, Jane Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 15PH-0346
FILING DATE: MARCH 100:
NAME: Licate, Jane Massey
REGISTRATION NUMBER: 15PH-0346
FILING APPLICATION NUMBER: 15PH-0346
FILING AP
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MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION 1.
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
TELEBRENCE/DOCKET NUMBER: 1SPH-0346
TELECOMMUNICATION INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               609-810-1454
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US-09-280-805-257
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          US-09-280-805-260/C
US-09-806-805-260/C
US-09-804-805-260/C
Patent No. 6184212
GENERAL INFORMATION:
PAPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. ANTION: EXRRESION
NUMBER OF SEQUENCES:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
COUNTRY: U.S.A.
ZIP: 08053
COUNTRY: U.S.A.
COUNTRY: U.S.A.
COUNTRY: INP COUNTRY: U.S.A.
COUNTRY: LISM PC
COUNTRY: INP COUNTRY: U.S.A.
COUNTRY: LING DATE: Herewith
CLASSIPICATION NUMBER: US/09/280,805
FILING DATE: Herewith
CLASSIPICATION NUMBER: US/09/280,805
FILING DATE: Herewith
CLASSIPICATION NUMBER: 132,27
REFERENCE/DOCKET NUMBER: 132,27
REFERENCE/DOCKET NUMBER: 132,27
RETERENCE/DOCKET NUMBER: 132,27
RETERENCE/DO
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Sequence 261, Application US/09280805
Fatent No. 6184212
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.38+02;
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COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Pred. ....
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Matches 20; Conservative
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COUNTRY: U.S.A.
7TP: 08053
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US-09-280-805-260/c
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APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Mariton
STATE: NJ
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MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: DISKETTE, 0.0

COMPUTER: DISKETTE, 0.0

COMPUTER: DISKETTE, 0.0

COMPUTER: DISKETTE, 0.0

COMPUTER: DATA:

APPLICATION NUMBER: US/09/280,805

FILING DATE: herewith
CLASSIFICATION NUMBER: 09/048,810

FILING DATE: March 26, 1998

ATTORNEY/AGENT INFORMATION:

NAME: Licata, Jane Massey

REGISTRATION NUMBER: 32,257

REGISTRATION NUMBER: 1599-0346

TELECOMMUNICATION INFORMATION:

TELEFAX: 609-810-1515

TELEFAX: 609-810-1515

SEQUENCE CHARACTERISTICS:

LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.8%; Score 20; DB 1; L
100.0%; Pred. No. 3.3e+02;
                                             APPLICATION ATTE: herewith CLASSIPICATION: DATE: herewith CLASSIPICATION DATA: APPLICATION DATA: APPLICATION DATE: APPLICATION NUMBER: 09/048,810 FILING DATE: MARCH 26,1998 ATTORNEY/AGENT INPORMATION: NAME: Licata, Jane Massey, REGISTRATION NUMBER: 32,257 REFERENCE/DOCKET NUMBER: 1SPH-0346; TELEPHONE: 609-810-14515; TELEPHONE: 609-810-14515; TELEPHONE: 609-810-14515; TELEPHONE: CHARACTERISTICS: LENGTH: 20 base pairs: LENGTH: 20 base pairs: LENGTH: 20 base pairs: TYPE: Nucleic Acid STRANDEDNESS: Single STRANDEDNESS: Single STRANDEDNESS: Single STRANDESE: Yes US-09-280-805-261
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity 100.0%; Pred. No. 3...
Matches 20; Conservative 0; Mismatches
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
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Patent No. 6184212
GENERAL INFORMATION:
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SENERAL INFORMATION:
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                                                                                                                                                                                                                                                  Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
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; Mismatches 0; Indels
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ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
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OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIPICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26,1998
ATTOCNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
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                                                                                                                                                                                                                                                                                                                                                                                                                   2298 GGTCTCGATCTCCTGACCTC 2317
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           20 GGTCTCGATCTCCTGACCTC 1
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INFORMATION FOR SEQ ID NO: 20 SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
; TYPE: Nucleic Acid
STRANDEDNESS: Single
; TOPOLOGY: Linear
; ANTL SENSE: Yes
US-09-280-805-262
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-280-805-263
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APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCES. 271
ADDRESSEE: Law Offices of Jane Massey Licata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    '18-09-280-805-265/c
'Sequence 265, Application US/09280805
'Sequence 265, Application US/09280805
'Patent No. 6184212
'GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
'ITILE OF INVENTION: EXPRESSION
'TITLE OF INVENTION: EXPRESSION
'NUMBER OF SEQUENCES: 271
'CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION NUMBER: 09/048,810
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
PRIOR APPLICATION NUMBER: 32,199
ATTORNEY/CART INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 15PH-0346
TELECOMMUNICATION INFORMATION:
TELEBHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                   ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 Bast Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2319 TGATCCGCCCACCTCGGCCT 2338
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         20 TGATCCGCCCACCTCGGCCT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 609-610-1454
INFORMATION FOR SEQ ID NO: 22
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 100.0
Matches 20; Conservative
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MEDIUM TYPE: DISKETT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: Linear
TOPOLOGY: Linear
ANTI-SENSE: Yes
US-09-280-805-264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM PC
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTRY: U.S.A.
ZIP: 08053
                                                                                                                                                                                                                                                     U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Marlton
                                                                                                                                                                                                                         STATE: NJ
COUNTRY: U.
ZIP: 08053
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US-09-280-805-267/c

US-09-280-805-267/c

is Sequence 267, Application US/09280805

is Patent No. 6184212

is GENERAL INFORMATION:

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.

APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: ANTISENSE MODULATION OF HUMAN MDM2

TITLE OF INVENTION: EXPRESSION

NUMBER OF SEQUENCES: 27

CORRESPONDENCE ADDRESS:

ADDRESSE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street

CITY: Marlton

STREET: NJ
                                                                                  Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 20;
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MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
OPERATING SYSTEM: MINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION WHOBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA: APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 15PH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-280-805-268/c
; Sequence 268, Application US/09280805
; Patent No. 6184212
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1; Le
                                                                               0.8%; Score 20; DB 1; L. 100.0%; Pred. No. 3.3e+02;
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                                                                                                      Best Local Similarity 100.0%; Pred. No. 3...
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                   2334 GGCCTCCCAAAGTGCTGGGA 2353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2341 CAAAGTGCTGGGATTACAGG 2360
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A.
          Yea
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       ; ANTI-SENSE:
US-09-280-805-266
                                                                                    Query Match
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APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 271
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.38+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:

MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
CURRENT APPLICATION DATA:

PILING DATE: herewith
CLASSIFICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTORNEY AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE COOPERT NUMBER: 15.PH-0346
TELECOMMUNICATION INFORMATION:
TELEPRAN: 609-810-1515
TELEPRAN: 609-810-1454
INFORMATION FOR SEQ ID NO: 266:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: NUCLALS ACID
COMPUTER: SINGLE
TYPE: NUCLALS ACID
COMPUTER: SINGLE
TYPE: NUCLALS ACID
COMPUTER: SINGLE
                                                                                                                                                                      NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION:
TELEPHONE: 609-810-1515
APPLICATION NUMBER: US/09/280,805
                    FILING DATE: herewith CLASSIFICATION:
PRIOR APPLICATION DATA: APPLICATION NUMBER: 09/048,810 FILING DATE: March 26, 1998 ATTORNEY/AGENT INFORMATION: NAME: Licata, Jane Massey
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2325 GCCCACCTCGGCCTCCCAAA 2344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 419
US-08-280-805-266/c
; Sequence 266, Application US/09280805
; Patent No. 6184212
                                                                                                                                                                                                                                                                                               TELEPAX: 609-810-1444

INFORMATION FOR SEQ ID NO: 265:
SEQUENCE CHARACTERISTICS:
LENGITH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
US-09-280-805-265
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 GCCCACCTCGGCCTCCCAAA 1
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Best Local Similarity 100.0
Matches 20; Conservative
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Gaps

Page 151

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37 GGCCCTGTGTGTCGGAAAGA 56
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         ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COUNTRY: U.S.A. ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ANTI-SENSE: Yes
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US-09-046-810-3/c
US-09-046-810-3/c
; Sequence 3, Application US/09048810
; Patent No. 6228921
; GENERAL INFORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE
; TITLE OF INVENTION: MODULATION OF HUMAN MDM2 EXPRESSION
; NUMBER OF SEQUENCES: 3
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
    ANTISENSE MODULATION OF HUMAN MDM2
EXPRESSION
271
                                                                                                                                                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER IBM PC
COMPUTER: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
COMPUTER: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM PC
COMPUTER: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: DISKETTE, 3.6 INDOME
SOFTWARE: WORDER: US/09/280,805
FILING DATE: Herewith
CLASSIFICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: MATCH 26, 1998
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REFERENCE/DOCKET NUMBER: 132,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELECHNONE: GOO-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
TITLE OF INVENTION: ANTISENSE MODULATION OF HUNDABER OF SEQUENCES: 271
NUMBER OF SEQUENCES: 271
NUMBER OF SEQUENCES: 271
ADDRESSED: Law Offices of Jane Massey Licata STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/048,810
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2351 GGATTACAGGCATGAGCCAC 2370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 268:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       20 GGATTACAGGCATGAGCCAC 1
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COUNTRY: U
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; Sequence 4, Application US/09048810
; Sequence 4, Application US/09048810
; Patent No. 6238921
; APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE
; TITLE OF INVENTION: MODULATION OF HUMAN MDM2 EXPRESSION
; VUMBER OF SEQUENCES: 32
; CORRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; STREET: 66 East Main Street
; CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match (1) 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READALE FORM:
MEDIUM TYER DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: WORDERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/048,810
FILING DATE: Herewith
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0302
TELECHOME: 609-779-2400
TELECHOME: 609-810-1454
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: CLARACTERISTICS:
LENGTH: CLARACTERISTICS:
NAME: Licata, Jane Massey

REGISTRATION UNDRER: 32,257

REFERENCE/DOCKET NUMBER: 15PH-0302

TELECOMMUNICATION INFORMATION:

TELEFONE: 609-779-2400

TELEFONE: 609-810-1454

INFORMATION FOR SEQ ID NO: 3:

SEQUENCE CHARACTRISTICS:

LENGTH: 20 base pairs

LENGTH: 20 base pairs

TYPE: Nucleic Acid

STRANDEDNESS: Single

STRANDEDNESS: Single

MANTI-SENSE: Yes

US-09-048-810-3
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0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE
TITLE OF INVENTION: MODULATION OF HUMAN MDM2 EXPRESSION
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/048,810
FILING DATE: herewith
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKT NUMBER: 1SPH-0302
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
Best Local Similarity 100.0%; Pred. No. 3.:
Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    147 ATTAGTGCGTACGAGCGCCC 166
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Patent No. 6238921
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; TYPE: Nucleic Acid
; STRANDEDNESS: Single
; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-048-810-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 609-810-1454
                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: Linear ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 08053
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                                                                                                                        Sequence 5, Application US/09048810
| Patent No. 6238921
| GENERAL INFORMATION:
| APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia
| TITLE OF INVENTION: MUDULATION OF HUMAN MDMZ EXPRESSION NUMBER OF SEQUENCES: 32
| CORRESPONDENCE ADDRESS: 32
| ADDRESSEE: Law Offices of Jane Massey Licata | CITY: Marlton | CITY: Marlton | CITY: Marlton | COUNTRY: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; iive 0; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: C.C....

ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: WORDDERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/048,810
FILING DATE: herewith
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 12,257
REFERENCE/DOCKET NUMBER: 12,257
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-779-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE OLICONUCLEOTIDE
TITLE OF INVENTION: MODULATION OF HUMAN MDM2 EXPRESSION
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Mailton
STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
COMPITER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GCTG 114
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                  20 GGCCCTGTGTGTCGGAAGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE: 609-779-2400
TELEPRAT: 609-810-1454
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs Type: Nucleic Acid STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity 100.
1es 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   95 CTCTGACCGAGATCCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Linear
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; ANTI-SENSE: Yes
US-09-048-810-5
                                                                                                      US-09-048-810-5/c
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66 East Main Street
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ANTI-SENSE: Yes
                                                                  U.S.A.
                         Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-048-810-9
                                                                       COUNTRY:
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Sequence 8, Application US/09048810

Fatent No. 6238921

GENERAL INFORMATION:

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.

CORRESPONDENCES: 32

CORRESPONDENCE ADDRESS:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street

CITY: Marlton

STREET: NJ

COUNTRY: U.S.A.

ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                    Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9, Application US/09048810
Patent No. 6238921
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Breet P. Monia
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE
TITLE OF INVENTION: MODULATION OF HUMAN MDM2 EXPRESSION
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
                                                                  0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
                 DB 1; Le. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: LIBM 486
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/048,810
FILING DATE: herewith
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 32,257
REGISTRATION NUMBER: 15PH-0302
TELECOMMUNICATION NUMBER: 15PH-0302
TELECOMMUNICATION NUMBER: 15PH-0302
TELECOMMUNICATION NUMBER: 15PH-0302
TELEPHONE: 609-910-1454
INFORMATION FOR SEQ ID NO: 8:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
            Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.: Matches 20; Conservative 0; Mismatches
                                                                                                               181 GAGAGTGGAATGATCCCCGA 200
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US-09-048-810-9/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE OLIGONUCLECTIDE
TITLE OF INVENTION: MODULATION OF HUMAN MDM2 EXPRESSION
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COUNTRY: 0.0....
ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
PLICATION NUMBER: US/09/048,810
FILING DATE: herewith
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REFERENCE/DOCKET NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0302
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match

0.8%; Score 20; DB 1; Lu
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
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                                                                     COMPUTER: 1BM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/048,810
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-048-810-10/c
; Sequence 10, Application US/09048810
Patent No. 6738921
; GENERAL INFORMATION:
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NAME: Licata, Jane Massey
REGISTATION UNDRER: 32.257
REFERENCE/DOCKET NUMBER: 1SPH-
TELECOMMUNICATION INFORMATION:
TELEFRAM: 609-779-2400
TELEFRAM: 609-779-2400
TELEFRAM: 209-779-2400
TELEFRAM: 20 base pairs
TENERAL: 20 base pairs
TYPE: NUCleic Acid
STRANDENNESS: Single
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                                                                                                                                                                             0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 11, Application US/09048810
Patent No. 6238921
GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE
TITLE OF INVENTION: MODULATION OF HUMAN MDM2 EXPRESSION
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COUNTY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
COMPUTER: IBM 486
COMPUTER: SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/048,810
FILLING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Law Offices of Jane Massey Licata STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ISPH-0302
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US-09-048-810-12/c
; Sequence 12, Application US/09048810
; Patent No. 6238921
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     331 CTGTACCTACTGATGGTGCT 350
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
CLASSIFICATION:
ATTORNEY/ACENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH
TELECOMMUNICATION INFORMATION:
TELECHONE: 609-779-2400
TELEPAX: 609-810-1454
                                                                                                                                                                                                                                                                                              20 AGCAGGCAAATGTGCAATAC 1
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INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                  ; LENGTH: 20 base pairs
; TYPE: Nucleic Acid
; STRANDEDNESS: Single
; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-048-810-10
                                                                                                                                                                                                    Best_Local Similarity 100.0
Matches 20; Conservative
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Best Local Similarity 100.
Matches 20; Conservative
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: Linear

ANTI-SENSE: Yes

US-09-048-810-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                               RESULT 430
US-09-048-810-11/c
                                                                                                                                                                               Query Match
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Sequence 13, Application US/09048810

Sequence 13, Application US/09048810

GENERAL INCORMATION:

APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION:

NUMBER OF SEQUENCES: 32

CORRESPONDENCE ADDRESS:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street

CITY: Mailton

CITY: Mailton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 20; DB 1; Length 20;
Loren J. Miraglia, Pamela Nero, Mark J.
Graham, Brett P. Monia
VENTION: ANTISENSE OLIGONUCLEOTIDE
VENTION: MODULATION OF HUMAN MDM2 EXPRESSION
                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: LBM 486
COMPUTER: WONDER: CBM
SOFTWARE: WONDER: CBM
CLASSIP CATION NUMBER: US/09/048,810
FILING DATE: herewith
CLASSIP CATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
CLASSIP CATION NUMBER: 159H-0302
FRETERROVE/DOCKET NUMBER: 159H-0302
TELLEPHONE: 609-779-2400
TELLEPHONE: 609-779-2400
TELLEPRAN: 609-810-1454
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
TENMATH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        3.3e+02;
                         APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE
TITLE OF INVENTION: MODULATION OF HUMAN MDM2 EN
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               100.0%; Pred. ...
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CURRENT APPLICATION DATA:
AAPPLICATION NUMBER: US/09/048,810
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         617 GATCTACAGGAACTTGGTAG 636
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 08053
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: Sing
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Best Local Similarity
Matches 20; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            U.S.A.
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                                                                                                                                                                                                                                    COUNTRY: U
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY:
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Best Loca
Matches
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                                                                                                                                                                                                                                                                               Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTIGENSE OLIGONUCLEOTIDE
TITLE OF INVENTION: MODULATION OF HUMAN MDM2 EXPRESSION
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
                                                                                                                                                                                                                                                                             Score 20; DB 1; Le
Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Pred. No. ...
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/048,810
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                         Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.: Matches 20; Conservative 0; Mismatches
                                       ISPH-0302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ISPH-0302
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-048-810-14/c; Sequence 14, Application US/09048810 Patent No. 6238921; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION UNDRER: 32,25F
REFERENCE/DOCKET UNDRER: ISPH-
TELECOMMUNICATION INFORMATION:
TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                    20 AGTGTAGAATTTGAAGTTGA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          20 rrgargriccrgarrgraaa 1
                                                                                      TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: IS
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-779-2400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
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Best Local Similarity
Matches 20; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Linear
Yes
                                                                                                                                                                                                 ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-048-810-13
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; Sequence 16, Application US/09048810
; Patent No. 6238921
; GENERAL INFORMATION:
; APPLICANT: Loraham, Brett P. Monia
; TITLE OF INVENTION:
; TITLE OF INVENTION: MODILATION OF HUMAN MDM2 EXPRESSION
; NUMBER OF SEQUENCES: 32
; CORRESPONDENCE ADDRESS:
; ADDRESSES: Law Offices of Jane Massey Licata
; STREET:, 66 East Main Street
; CITY: Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.8%; Score 20; DB 1; Length 20;
100.0%; Pred. No. 3.38+02;
iive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                EXPRESSION
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ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: WORDPERECT 5.1
CURRENT APPLICATION DATA:
PILING DATE: herewith
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REFERENCE/DOCKET NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0302
TELECOMMUNICATION INFORMATION:
                       Sequence 15, Application US/09048810
| Patent No. 6238921
| GENERAL INFORMATION:
| APPLICANT: Graham, Brett P. Monia | APPLICANT: Graham, Brett P. Monia | TITLE OF INVENTION: MODULATION OF HUMAN MDM2 EXPRESSIONUMBER OF SEQUENCES: 32 CORRESPONDENCE ADDRESS: ADDRESSEE: Law Offices of Jane Massey Licata | STREET: Marlton | Main Street | CITY: Marlton | Main Street | Marlton | Marlton | Main Street | Marlton | Marlto
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM 486
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1695 TTTACATGTGCAAAGAAGCT 1714
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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Best Local Similarity
Matches 20; Conserva
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US-09-048-810-15/C
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SEQUENCE CHARACTERICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
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                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-048-810-19/c
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Patent No. 6238921

GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE
TITLE OF INVENTION: MODULATION OF HUMAN MDM2 EXPRESSION
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 20;
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COMPUTRY: NJ

ZIP: 08053
COMPUTRY: READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/048,810
FILING DATE: herewith
CLASSIFICATION NUMBER: US/09/048,810
FILING DATE: harewith
CLASSIFICATION NUMBER: 32.257
REFERENCE/DOCKET NUMBER: 32.257
REFERENCE/DOCKET NUMBER: 1SPH-0302
TELECHONE: 609-810-1454
INFORMATION FOR SEQ ID NO: 17: SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
TOORAGE: Yes

ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Sco. 100.0%; Pred. No. ...
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS SOFTWARE: WORDPERFECT 5.1
                                                                                                                                    NAME: Licata, Jane Massey
REGISTRATION UNWBER: 32,257
REFERENCE/DOCKET NUMBER: 15PH-0302
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-79-2400
TELEPHONE: 609-810-1454
INFORMATION FOR SEQ ID NO: 16: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                           US/09/048,810
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                                                                              FILING DATE: herewith
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.8*
Best Local Similarity 100.0
Matches 20; Conservative
                             CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
FILING DATE: herewith
                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-048-810-16
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Gaps
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Sequence 18, Application US/09048810

Sequence 18, Application US/09048810

Patent No. 6238921

APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J. APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE

TITLE OF INVENTION: MODULATION OF HUMAN MDM2 EXPRESSION

NUMBER OF SEQUENCES: 32

CORRESPONDENCE ADDRESS:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 1.9, Application US/09048810
Patent No. 6238921
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE
TITLE OF INVENTION: ANDULATION OF HUMAN MDM2 EXPRESSION
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
  Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTE: DESCRIPE, 3.5 INCH, 1.44 MD STORAGE MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE COMPUTER: IBM 486 COMPUTER: BM 486 COPERATING SYSTEM: WINDOWS FOR WORKGROUPS SOFTWARE: WORDPERFECT 5.1 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/048,810 FILLNG DATE: herewith CLASSIFICATION:
Query Match
0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No., 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 20; DB 1; L; Pred. No. 3.3e+02; 0; Mismatches 0:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFRENCE/DOCKET NUMBER: ISPH-0302
TELECOMMUNICATION INFORMATION:
TELEFANE: 609-779-2400
TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 18:
                                                                                                           1785 TAGTTGACCTGTCTATAAGA 1804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1818 CTAACTATATAACCCTAGGA 1837
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Best Local Similarity 100.0%;
Matches 20; Conservative 0
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US-09-048-810-21/c

Sequence 21, Application US/09048810

Patent No. 6238921

GENERAL INFORMATION:

APPLICANT: Graham, Brett P. Monia

TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE

TITLE OF INVENTION: MODULATION OF HUMAN MDM2 EXPRESSION

NUMBER OF SEQUENCES: 32

CORRESPONDENCE ADDRESS:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street

CITY: Mailton

STATE: NJ
                                                                                                                                                0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MD STORAGE
COMPUTER: IBM 486
COMPUTER: IBM 486
COPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: WORDERFECT 5.1
CURRENT APPLICATION NATA:
APPLICATION NUMBER: US/09/048,810
FILING DATE: Herewith
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0302
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECHORIES 609-779-2400
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; Sequence 22, Application US/09048810
; Patent No. 6238911
; GENERAL INPORMATION:
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0.8%; Score 20; DB 1; Lo
100.0%; Pred. No. 3.3e+02;
tive 0; Mismatches 0;
                                                                                                                                                             100.0%; Prec. ...
                                                                                                                                                                                                                                   2132 AGTGCAGTGGGTGATCTTGG 2151
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 AGTCATCTGCCACCACACCT
                                                                                                                                                                                          20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 20 base pairs
TYPE: Nucleic Acid
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Best Local Similarity 100.
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
        20 base pairs
                                                : Single
                         TYPE: Nucleic Acid
STRANDEDNESS: Sing
                                                                                                                                                                     Best Local Similarity
                                                                   Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; ANTI-SENSE: Yes
US-09-048-810-21
                                                                   TOPOLOGY:
                                                                                    ; ANTI-SENSE:
US-09-048-810-20
      LENGTH:
                                                                                                                                                  Query Match
                                                                                                                                                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: LOTEN J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTIENSE OLIGONUCLEOTIDE
TITLE OF INVENTION: MODULATION OF HUMAN MDM2 EXPRESSION
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: WORDPERRECT 5.1
CURRENT APPLICATION DATA:
RILING DATE: herewith
                                                         Score 20; DB 1; Le
Pred. No. 3.3e+02;
                                                                                                                      COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/048,810
FILING DATE: herewith
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.8%; Score 20; DB Best Local Similarity 100.0%; Pred. No. 3.: Matches 20; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                ISPH-0302
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-048-810-20/c; Sequence 20, Application US/09048810 Patent No. 6238921; GENERAL INFORMATION:
                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUBBER: 32,257
REFERENCE/DOCKET NUMBER: 15PH-
TELECOMMUNICATION INFORMATION:
TELEFAX: 609-179-2400
TELEFAX: 609-180-1454
INFORMATION FOR SEQ ID NO: 19: SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
TYPE: Nucleic Acid
TYPE: Nucleic Acid
TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
NAME: Licate, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-779-2400
TELEPAK: 609-810-1454
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Linear
Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A.
                                          U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
  Marlton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
ANTI-SENSE:
CITY: Ma:
STATE: N.
COUNTRY:
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Query Match
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APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTIESENSE OLIGONUCLEOTIDE
TITLE OF INVENTION: MODULATION OF HUMAN MDM2 EXPRESSION
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
STATE: Marlton
STATE: 08053.
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE
TITLE OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1; Length 20; 3.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                             COMPUTER: NU
COMPUTER: NU
COMPUTER: NU
COMPUTER: D8053
COMPUTER: D15KETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: D16M AND AGE
SOFTWARE: WORDERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/048,810
FLLIG DATE: NEEWAITON:
ATTORNEY/AGENT INFORMATION:
NAME: Licata, Jane Massey
REFERENCE/DOCKET NUMBER: 1257
TELECHONE: 609-779-7400
TELECHONE: 609-779-7400
TELECHONE: 609-779-7400
TELECHONE: 609-779-7400
TELECHONE: O59-779-7400
TTYPE: NUCLEIC Acid
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MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/048,810
FILING DATE: herewith
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.8%; Score 20; DB 100.0%; Pred. No. 3.3
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; Sequence 25, Application US/09048810
; Patent No. 6238921
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.8%
Best Local Similarity 100.0
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-09-048-810-22
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APPLICANT: Brett P. Monia
APPLICANT: Lex M. Cowsert
TITLE OF TIVENTION: ANTISENSE MODULATION OF TELOMERIC REPEAT BINDING FACTOR 2 EXPRES
FILE REFRENCE: RTS-0106
CURRENT APPLICATION NUMBER: US/09/467,642
CURRENT FILING DATE: 1999-12-20
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 71
LENGTH: 20
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APPLICANT: Brett P. Monia
APPLICANT: Brett P. Monia
APPLICANT: Lex M. Coweert
TITLE OF INVENTION: ANTISENSE MODULATION OF TELOMERIC REPEAT BINDING FACTOR 2 EXPRES
FILE REFERENCE: RTS-0106
CURRENT APPLICATION NUMBER: US/09/467,642
CURRENT APPLICATION NUMBER: 1999-12-20
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 73
LENGTH: 20
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100.0%; Pred. No. 3.3e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                    0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; FEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-09-467-642-71
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ORGANISM: Artificial Sequence
FEATURE:
JOTHER INFORMATION: Antisense Oligonucleotide
US-09-467-642-73
                                                                                                                                                                                                                                                                                                                                            Query Match 0.8%; Score zv; LD Best Local Similarity 100.0%; Pred. No. 3.3 Matches 20; Conservative 0; Mismatches
                 REFERENCE/DOCKET NUMBER:
TELECOMUNICATION INFORMATION:
TELEPHONE: 609-779-2400
TELEPHOR: 609-719-2400
TELEPHOR: 609-719-2400
TELEPHOR: 609-810-1454
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDENNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.8%; Score 20;
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; Sequence 73, Application US/09467642
; Patent No. 6300132
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 71, Application US/09467642
Patent No. 6300132
GENERAL INFORMATION:
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REGISTRATION NUMBER: 32,257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-467-642-71/c
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US-09-048-810-25
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Pred. No. 3.3e+02;

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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/060,299
FILING DATE: 15-APR-1998
CLASSIFICATION: 435
PRICK APPLICATION DATA:
APPLICATION NUMBER: US 60/043,553
FILING DATE: 15-APR-1997
PRICK APPLICATION DATA:
APPLICATION NUMBER: US 60/048,740
FILING DATE: 05-UIN-1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                              Sequence 3, Application US/09607529
Sequence 3, Application US/09607529
GENERAL INFORMATION:
APPLICANT: David Traver:
APPLICANT: David Traver:
APPLICANT: Kolch! Akashi
TITLE OF INVENTION: WHERENCE:
TITLE OF INVENTION: SUBSETS
FILE REPREBRUCE: STAN-126
CURRENT APPLICATION NUMBER: US/09/607,529
CURRENT FILING DATE: 1099-06-29
PRIOR APPLICATION NUMBER: 60/141,421
PRIOR APPLICATION NUMBER: 61/141,421
NUMBER OF SEQ ID NOS: 6
SOFTWARE: FASESEQ for Windows Version 4.0
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ADDRESSEE: Nixon and Vanderhye
STREET: 1100 No. 6545137th Glebe Road, Eighth Floor
CITY: Arlington
STATE: Virginia
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Todd, John A
APPLICANT: Hess, John W
APPLICANT: Gaskey, Charles T
APPLICANT: Cox, Roger D
APPLICANT: Gerhold, David
APPLICANT: Hey, Patricia
APPLICANT: Kawaguchi, Yoshihiko
APPLICANT: Merriman, Tony R
APPLICANT: Merziman, Tony R
APPLICANT: Merzker, Michael L
TITLE OF INVENTION: No. 6545137el Receptor
NUMBER OF SEQUENCES: 455
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              . Sequence 257, Application US/09060299; Patent No. 6545137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2344 AGTGCTGGGATTACAGGCAT 2363
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 AGTGCTGGGATTACAGGCAT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Todd,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-060-299-257/c
                                                                                        RESULT 447
US-09-607-529-3/c
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LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                          Sequence 62, Application US/0948856A
Patent No. 6316259
GENERAL INFORMATION:
APPLICANT: Breet P. Monia
APPLICANT: Breet P. Monia
APPLICANT: Breet M. Butler
APPLICANT: Madeline M. Butler
APPLICANT: Madeline M. Butler
APPLICANT: Madeline M. Butler
APPLICANT: APPLICA
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                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Agrawal, Sudhir
APPLICANT: Kandimalla, Ekambar R.
TITLE OF INVENTION: Peeudo-Cyclic Oligonucleobases
FILE REPERSONCE: 99,128-B
CURRENT APPLICATION NUMBER: US/09/540,699
CURRENT FILING DATE: 2000-03-31
PRIOR APPLICATION NUMBER: US 60/127,138
PRIOR APPLICATION NUMBER: US 60/174,642
PRIOR FILING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 26
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO S.2
FEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-09-488-856A-62
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                                                                                                                     2346 IGCIGGGATTACAGGCATGA 2365
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Sequence 2.5, Application US/09540699
Patent No. 6383752
GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
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                 Best Local Similarity 100. Matches 20; Conservative
                                                                                                                                                                                                                                                                                        RESULT 445
US-09-488-856A-62
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TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 257:
STRANDEDNESS: single
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                                                           US-09-402-923A-257
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COMPUTER: INDEPTY LINE
COMPUTER: INDEPTY LINE
COMPACTION SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATORIN Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/402,923A
FILING DATE: 14-F6b-2001
FILING DATE: 15-APR-1998
APPLICATION NUMBER: PCT/GB98/01102
FILING DATE: 15-APR-1997
APPLICATION NUMBER: US 60/043,553
FILING DATE: 15-APR-1997
APPLICATION NUMBER: US 60/048,740
FILING DATE: 05-JUN-1997
APTORNEY/AGENT INFORMATION:
NAME: J. SAGOÉF
                                                                                                                                                                                                                                                                                                                                     ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE: Nixon and Vanderhye
STREET: 1100 No. 6555654th Glebe Road, Eighth Floor
STREET: Arington
STATE: Virginia
COUNTRY: US
ZONTER: READBLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                         0.8%; Score 20; DB 1; Length 20; 100.0%; Pred. No. 3.3e+02; tive 0; Mismatches 0; Indels
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Kawaguchi, Yoshihiko
Merriman, Tony R
Metzker, Michael L
TITLE OF INVENTION: No. 6555654el LDL-Receptor
NUMBER OF SEQUENCES: 455
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 36,663
REFERENCE/DOCKET NUMBER: 620-81
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-402-923A-257/c
; Sequence 257, Application US/09402923A
; Patent No. 6555654
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                          2344 AGTGCTGGGATTACAGGCAT 2363
    REGISTRATION NUMBER: 36,663
REGISTRATION NUMBER: 620-3
TELECOWNUICATION INFORMATION:
TELEPHONE: (703)816-4091
TELEPHONE: (703)816-4100
INFORMATION FOR SEQ ID NO: 257:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single

US-09-060-299-257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: Todd, John A
Hess, John W
Caskey, Charles T
                                                                                                                                                                                                                                                                                                                                                                                                     20 AGTGCTGGGATTACAGGCAT 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Cox, Roger D
Gerhold, David
Hammond, Holly
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SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
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Best Local Similarity 100.0%
Matches 20; Conservative
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GENERAL INCORMATION:
APPLICANT: Mancharan, Muthiah
APPLICANT: GUZaev, Andrei F.
TITLE OF INVENTION: Process For Preparing Peptide Derivatized Oligomeric Compounds
FILE REFERENCE: ISIS4501
CURRENT APPLICATION NUMBER: US/09/658,517C
CURRENT FILING DATE: 2000-09-08
SOFTWARR: Patentin version 3.1
SSQ ID NO 2.2
LENGTH: 20
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APPLICANT: CHERIF, Dorra
APPLICANT: CHERIF, Dorra
TITLE OF INVENTION: FLUORESCENT PROBES FOR CHROMOSOME PAINTING
FILE REPERENCE: GENERT.06594US
CURRENT APPLICATION NUMBER: US/09/418,804A
CURRENT APPLICATION NUMBER: F 98/12957
EARLIER APPLICATION NUMBER: F 98/12957
EARLIER FILING DATE: 1998-10-15
SEQ ID NO 1
LENGTH: 20
Length 20;
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DB 1; Le
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0.8%; Score 20; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 3.3e+02;
Matches 20; Conservative 0; Mismatches 0;
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Query Match

0.8%; Score 20; DB
Best Local Similarity 100.0%; Pred. No. 3.3
Matches 20; Conservative 0; Mismatches
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                                                                                           2344 AGTGCTGGGATTACAGGCAT 2363
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US-09-658-517C-2/c
; Sequence 2, Application US/09658517C
; Patent No. 6559279.
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; OTHER INFORMATION: primer PCR Alu
US-09-418-804-1
                                                                                                                             20 AGTGCTGGGATTACAGGCAT 1
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Wed Jan 26 12:47:30 2005

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US-09-949-474A-2/C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     OTHER INFORMATION: Description of Artificial Sequence: No. 6656730el OTHER INFORMATION: Sequence
                                                          US-09-679-2994-70

US-09-679-2994-70

US-09-679-2994-70

Sequence 70, Application US/09679299A

Patent No. 6566135

GRNERAL INFORMATION:

APPLICANT: Workine L. Brown-Driver

APPLICANT: Hong Zhang

APPLICANT: Andrew T. Watt

TITLE OF INVENTION ANTIENSE MODULATION OF CASPASE 6 EXPRESSION

FILE REFERENCE: RTS-0187

CURRENT APPLICATION NUMBER: US/09/679,299A

CURRENT FILING DATE: 2000-10-04

NUMBER OF SEQ ID NOS: 164

SEQ ID NO 70

LENGTH: 20
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3.3e+02;
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GENERAL INFORMATION:
APPLICANT: Manoharan, Muthiah
TITLE OF INVENTION: Drug-Conjugated Oligomeric Compounds
FILE REFERENCE: ISIS3758
CURRENT APPLICATION NUMBER: US/09/334,130
CURRENT PLING DATE: 1999-06-15
NUMBER OF SEQ ID NOS: 9
SOFTWARE: Patentin Ver. 2.1
LENGTH: 20
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Patent No. 6/61883
GENERAL INFORMATION:
APPLICANT: Weissman, Irving L.
APPLICANT: Traver, David Jeffrey
APPLICANT: ARashi, Koichi
TITLE OF INVENTION: MAMMALIAN WYELOID PROGENITOR CELL
TITLE OF INVENTION: SUBSETS
                                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: Antisense Oligonucleotide US-09-679-299A-70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.8%; Score 20; DB
100.0%; Pred. No. 3.3
tive 0; Mismatches
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  20 CCCAGGCTGGAGTGCAGTGG
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Best Local Similarity 100.0
Matches 20; Conservative
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US-09-334-130-7/c
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Sequence 2, Application US/09949474A

Sequence 2, Application US/09949474A

Patent No. 676231

GENERAL INFORMATION:
APPLICANT: Manoharan, Muthiah
APPLICANT: Guzaev, Andrei P.
TITLE OF INVENTION: Process for Preparing Peptide Derivatized Oligomeric Compounds
FILE REFERENCE: ISIS4850
CURRENT APPLICATION NUMBER: US/09/949,474A

CURRENT APPLICATION NUMBER: 09/658,517
PRIOR FILING DATE: 2000-09-08
NUMBER OF SEQ ID NOS: 24
SEQ ID NO 2
IMPRORMED PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PLING DATE: 2000-09-08
SEQ ID NO 2
IMPRORMED PRIOR PRIO
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US-09-594-387-7
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APPLICANT: Mancharan, Muthiah

TITLE OF INVENTION: Ligand-Conjugated Oligomeric Compounds

TITLE OF INVENTION: Ligand-Conjugated Oligomeric Compounds

FILER REFERENCE: 15184390

CURRENT APPLICATION NUMBER: US/09/594,387

CURRENT FILING DATE: 2000-06-15

PRIOR FILING DATE: 1999-06-15

NUMBER OF SEQ ID NOS: 10

SOFTWARE: Patentin Ver. 2.1

LENGTH: 20
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. 3.3e+02;
FILE REFERENCE: STAN126CIP
CURRENT APPLICATION NUMBER: US/09/956,279
CURRENT FILING DATE: 2001-09-17
FRIOR APPLICATION NUMBER: 09/607,529
PRIOR FILING DATE: 2000-06-29
PRIOR FILING DATE: 1999-06-29
NUMBER OF SEQ ID NOS: 6
SOFTWARE: PastSEQ for Windows Version 4.0
SERIOTH: 20
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Homo sapiens
US-09-956-279-3
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Best Local Similarity
Matches 20; Conserva
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US-09-480-718-26
; Sequence 26, Application US/09480718
; Patent No. 6407062
; Patent No. 6407062
; Patent No. 6407062
; APPLICANT: Sherr, Charles J
; APPLICANT: Quelle, Dawn E
; APPLICANT: Quelle, Dawn E
; APPLICANT: Roussel, Martine F.
; APPLICANT: Prederique, Zindy
; TITLE OF INVERTION: ARF-19, A NOVEL REGULATOR OF THE MAMMALIAN CELL CYCLE
; TITLE OF INVERTION: ARF-19, A NOVEL REGULATOR OF THE MAMMALIAN CELL CYCLE
; CURRENT APPLICATION NUMBER: US/09/480,718
; CURRENT PILING DATE: 1998-08-06
; RARLIER PILING DATE: 1998-08-06
; SARLIER PILING DATE: 1998-08-06
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NOS: 48
; SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:

APPLICANT: Kent D. Taylor (Inventor)

APPLICANT: Maren T. Scheuner (Inventor)

APPLICANT: Jerome I. Scheuner (Inventor)

APPLICANT: Huiying Yang (Inventor)

TITLE OF INVENTION: Genetic Test to Determine

TITLE OF INVENTION: No. 6297014-responsiveness to Statin Drug Treatment

FILE REFERENCE: PO7 41878

CURRENT APPLICATION NUMBER: US/09/347,114A

CURRENT APPLICATION NUMBER: 1999-07-02

NUMBER OF SEQ ID NOS: 110

SOFTWARE: FastSEQ for Windows Version 4.0

SEQ ID NO 33

LENGTH: 26

TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Description of Artificial Sequence: Primer(sense)
US-09-480-718-26
     OTHER INFORMATION: /note= "These bases are listed 3'->5' left to
                                                                                                                                                                                                                                                                                                                                                .;
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                       OTHER INFORMATION: right...

NAME/KEY: misc feature

LOCATION: (20)...(21)

OTHER INFORMATION: /note= "3'-3' internucleotide linkage"

NAME/KEY: misc_feature

LOCATION: (8)

COTHER INFORMATION: /note= "n represents 5-DABCYL-thymidine"

US-09-540-699-12
                                                                                                                                                                                                                                                                                         Length 26;
                                                                                                                                                                                                                                                                                                                                              1; Indels
                                                                                                                                                                                                                                                                                      Query Match

0.8%; Score 20; DB 1; 1
Best Local Similarity 95.2%; Pred. No. 3.1e+02;
Matches 20; Conservative 0; Mismatches 1.
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                                                                                                                                                                                                                                                                                                                                                                                                     674 TGTGAGTGAGAACAGGTGTCA 694
                                                                                                                                                                                                                                                                                                                                                                                                                                                       21 TGTGAGTGAGAACNGGTGTCA 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA ORGANISM: Artificial Sequence
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Sequence 21, Application US/09949474A

Sequence 21, Application US/09949474A

Sequence 21, Application US/09949474A

GENERAL INFORMATION:

APPLICANT: Manohazan, Muthiah

APPLICANT: Manohazan, Andrei P.

TILLE OF INVENTION:

FILE REFERENCE: ISIS4850

CURRENT FILING DATE: 2001-09-07

FRIOR PPLICATION NUMBER: 09/658,517

PRIOR FILING DATE: 2000-09-08

NUMBER OF SEQ ID NOS: 24

SOUTHARE: Patentin version 3.2

SEQ ID NO 21

LENGTH: 20
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                                                                                                                                                                                                       Gaps
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                                                                                                                                      Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.3e+02; Matches 20; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.8%; Score 20; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.38+02; Matches 20; Conservative 0; Mismatches 0; Indels
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APPLICANT: Agrawal, Sudhir
APPLICANT: Agrawal, Sudhir
TILLE OF INVENTION: Pseudo-Cyclic Oligonucleobases; FILE REFREENCE: 99,128-B; CURRENT APPLICATION NUMBER: US/09/540,699; CURRENT APPLICATION NUMBER: US 60/127,138; PRIOR APPLICATION NUMBER: US 60/127,138; PRIOR FILING DATE: 1999-03-31; PRIOR FILING DATE: 2000-01-05; NUMBER OF SEQ ID NOS: 26; SOCTWARE: Patentin Ver: 2.0; SEQ ID NO 12.
                          ; FEATURE:
; OTHER INFORMATION: Synthetic construct
US-09-949-474A-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Synthetic construct US-09-949-474A-21
                                                                                                                                                                                                                                                      1695 TITACATGTGCAAAGAAGCT 1714
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ORGANISM: Artificial Sequence
PEATURE:
ORGANISM: Artificial Sequence
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COMPUTER READABLE FORM:
             GENERAL INFORMATION:
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                                                                                                                                                                                                                                          US-09-366-840-1
                                                                                                                                                                SEQ ID NO 1
LENGTH: 21
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                                                                                                                                                                                                                                                                        Sequence 7, Application US/09427104
Patent No. 6505652
GENERAL INFORMATION:
APPLICANT: Sherman M. Weissman
Jon J. Jonsson
TITLE OF INVENTION: Allele Frequency Differences Method for Phenotype
Cloning
                                                                                                       Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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                                                           Query Match 0.8%; Score 19.6; DB 1; Length 26; Best Local Similarity 84.6%; Pred. No. 3.2e+02; Matches 22; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 26;
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                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSEE: Dr. Sherman M. Weismann
Boyer Center for Molecular Medicine
Yale University School of Medicine
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 19.6; DB 1;
84.6%; Pred. No. 3.2e+02;
ative 0; Mismatches 4;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC
OPERATING SYSTEM: MS DOS
SOFTWARE: WOR'D PROCESSING
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/427,104
FILING DATE: 26-Oct-1999
PRIOR APPLICATION NUMBER: US 60/029,512
FILING DATE: 28-OCT-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: United States of America
ZIP: 06534-0812
COMPUTER READABLE PORM:
MEDIUM TYPE: 3.5" 1.44 Mb diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Mary M. Krinsky
REGISTRATION NUMBER: 32423
REFERENCE/DOCKET NUMBER: OCR-816
                                                                                                                                        2102 GACCGAGTCTTGCTCTGTTACCCCAGG 2127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2099 TGAGACCGAGTCTTGCTCTGTTACCC 2124
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DESCRIPTION: DNA
SEQUENCE DESCRIPTION: SEQ ID NO: 7:
                                                                                                                                                                            26 GACACAGICICGCICAGITACCCAGG 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: 203-773-9544
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 1, Application US/09366840 ; Patent No. 6228345
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 26 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear MOLECULE TYPE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 84.00,
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                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
; ORGANISM: Homo sapiens
US-09-347-114A-33
                                                                                                                                                                                                                                                        US-09-427-104-7/c
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                                                                                                                                                                                                                                                                                    Length 21;
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Patent No. 6337192

GENERAL INFORMATION:

APPLICANT: Bartel, Paul L.

APPLICANT: Tavtiglan, Sean V.

TITLE OF INVENTION: MMSC1 - An MMAC1 Interacting Protein

TITLE OF INVENTION: MMSC1 - An WAC1

CURRENT PLICATION NUMBER: US/09/233,086

CURRENT FILING DATE: 1999-01-19

EARLIER APPLICATION NUMBER: US 60/071,861

EARLIER PLING DATE: 1999-01-20

NUMBER OF SEQ ID NOS: 65

SOFTWARE: Patentin Ver. 2.0

SEQ ID NO 61

LENGTH: 21
                                                                                                                                                                                                                                                                                                                                1; Indels
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APPLICANT: OSSOWSKI, Liliana
TITLE OF INVENTION: In Vivo Assay for Intravasation
FILE REFERENCE: A32590 70165.0550
CURRENT APPLICATION WIMBER: US/09/366,840
CURRENT FILING DATE: 1999-08-04
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                    0.8%; Score 19.4; DB 1;
95.2%; Pred. No. 3.5e+02;
tive 0; Mismatches 1;
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Pred. No. 3.5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
APPLICANT: Hardy, John A.
APPLICANT: Goate, Alison M.
TITLE OF INVENTION: MUTANT $182 GENES
NUMBER OF SEQUENCES: 24
CORRESPONDENCE ADDRESS:
ADDRESSEB: SmithKline Beecham Corporation
STREET: 709 Swedeland Road
CITY: King of Prussia
STREET: DA
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Mismatches
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                              20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                        Best Local Similarity
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                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-233-086-61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 464
US-08-670-479-11
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Gaps
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Sequence 10, Application US/08629339

Sequence 10, Application US/08629339

GENERAL INFORMATION:

TITLE OF INVENTION: METHODS FOR DIAGNOSING AN INCREASED
TITLE OF INVENTION: METHODS FOR DIAGNOSING AN INCREASED
TITLE OF INVENTION: RISK OF BREAST OR OVARIAN CANCER
NUMBER OF SEQUENCE: 14

CORRESPONDENCE: 14

ADDRESSEE: SUGHRUE: MINN, ZINN, MACPEAK & SEAS
STREET: 2100 Fennsylvania Avenue, N.W., Suite 800
CITY: Washington, D.C.

COUNTRY: U.S.A.

COUNTRY: U.S.A.

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible

STATE: 18M PC Compatible

CORPUTER: Ploppy disk
COMPUTER: Patentin Release #1.0, Version #1.25

COMPUTER: PLOPEY WASHINGS SYSTEM: US/08/629,939
FILING DATE: 12-APRIL-1996

CURSSIFICATION NUMBER: US/08/629,939
FILING DATE: 12-APRIL-1996

CIRSEPHORE: (202) 293-7860

INFORMATION FOR SEQ ID NO: 10: SEQUENCE CHARACTERISTICS:

LENGTH: 19 base pairs

TYPE: nucleic acid

STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.8%; Score 19.2; DB 1; Length 24; Best Local Similarity 87.5%; Pred. No. 3.5e+02; Matches 21; Conservative 0; Mismatches 3; Indels
                  PILING DAID:

CLASSIFICATION 1435

PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/029,679
FILING DATE: 30-OCT-1996
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATE: US 60/028,056
FILING DATE: US -0.CCT-1996
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/025,719
FILING DATE: US -0.CCT-1996
ATTORNY/ACENT INFORMATION:
NAME: Wilson, Mark B.
FREGISTRATION NUMBER: ARCD:272
FREERENGE/DOCKET NUMBER: ARCD:272
FREERENGE/DOCKET NUMBER: ARCD:272
FILEDEMONE: 512/474-7577
INFORMATION FOR SEQ ID NO: 122:
SEQUENCE CHARACTERISTICS:
LENGRHOR TATA-7577
INFORMATION FOR SEQ ID NO: 122:
SEQUENCE CHARACTERISTICS:
LENGRHOR TATA-7577
INFORMATION FOR SEQ ID NO: 122:
SEQUENCE CHARACTERISTICS:
LENGRHOR TATA-7577
INFORMATION FOR SEQ ID NO: 122:
SEQUENCE CHARACTERISTICS:
LENGRHOR TATA-7577
INFORMATION FOR SEQ ID NO: 122:
SEGUENCE CHARACTERISTICS:
LENGRHOR TATA-7577
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LENGRHOR TATA-7577
INFORMATION FOR SEQ ID NO: 122:
SEGUENCE CHARACTERISTICS:
LENGRHOR TATA-7577
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SEGUENCE CHARACTERISTICS:
LENGRHOR TATA-7577
INFORMATION FOR SEQ ID NO: 122:
SEGUENCE CHARACTERISTICS:
LENGRHOR TATA-7577
INFORMATION FOR SEQ ID NO: 122:
SEGUENCE CHARACTERISTICS:
LENGRHOR TATA-7577
INFORMATION FOR SEQ ID NO: 122:
SEGUENCE CHARACTERISTICS:
LENGRHOR TATA-7577
INFORMATION FOR SEQ ID NO: 122:
SEGUENCE CHARACTERISTICS:
LENGRHOR TATA-7577
INFORMATION FOR SEQ ID NO: 122:
SEGUENCE CHARACTERISTICS:
SEGU
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2099 TGAGACCGAGICTIGCTCTGTTAC 2122
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           , TOPOLOGY:
US-08-927-219-122
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Sequence 123, Application US/08927219
Sequence 124, Aracya
APPLICANT: Vamagata, Kazuya
APPLICANT: Purtuta, Hioto
APPLICANT: Purtuta, Hioto
APPLICANT: Horitawa Yukio
APPLICANT: AND HNF-4ALPHA
AND HNF-4ALPHA
AND HNF-4ALPHA
CORRESPONDENCE: Arnold, White & Durkee
STREET: P.O. Box 4433
CITY: Houston
STREET: P.O. Box 4433
CITY: Houston
STRATE: Texas
COUNTRY: USA
ZIP: 77210
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: IBM PC Compatible
COMPUTER: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION NUMBER: US/08/927,219
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/670,479
FILING DATE: 26-UUN-1996
CLASSIFICATION NUMBER: 60/001,500
FILING DATE: 18-UL-1996
APPLICATION NUMBER: 60/001,800
FILING DATE: 18-UL-1996
APPLICATION NUMBER: 60/001,800
FILING DATE: 02-AUG-1955
ATTORNEY/AGENT INFORMATION:
NAMME: Han, William T
REGISTRATION NUMBER: 34,344
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           34,344
BER: P50361
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: PS
TELECOMMUNICATION INFORMATION:
TELEPHONE: 610-270-5219
TELEPAX: 610-270-5090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: CDNA
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FRAGMENT TYPE:
ORIGINAL SOURCE:
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US-08-759-873-10

Sequence 10, Application US/08759873

Sequence 10, Application US/08759873

Sequence 10, Application US/08759873

Sequence 10, Application US/08759873

TITLE OF INVENTION: METHODS FOR DIAGNOSING AN INCREASED RISK

TITLE OF INVENTION: OF RREAST OR OVARIAN CANCER

NUMBER OF SEQUENCES: 14

CORRESPONDENCE ADDRESS:

ADDRESSEE: SUGHRUE, MION, ZINN, MACPEAK & SEAS

STREET: 2100 Pennsylvania Avenue, N.W., Suite 800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Scheffler, Immo E. TITLE OF INVENTION: Mammalian Artificial Chromosomes and TITLE OF INVENTION: Methods of Using Same NUMBER OF SEQUENCES: 16
                                                                                             Length 19;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 19;
                                                                                                                                    0; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPOTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOGTWARE: PATENTIN Release #1.0, Version #1.25
SOGTWARENT APPLICATION DATA:
APPLICATION NUMBER: US/08/759,873
FILING DATE: 12-APRIL-1996
CLASSIPICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: KIT, Gordon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 19; DB 1; Le
100.0%; Pred. No. 3.8e+02;
tive 0; Mismatches 0;
                                                                                      Query Match 0.8%; Score 19; DB 1; L. Best Local Similarity 100.0%; Pred. No. 3.8e+02; Matches 19; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REGISTRATION NUMBER: 30,764
REFERENCE/DOCKET NUMBER: A-6612
TELECOMYUNICATION INFORMATION:
TELEPHONE: (202) 293-7060
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
                                                                                                                                                                             2342 AAAGTGCTGGGATTACAGG 2360
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2342 AAAGTGCTGGGATTACAGG 2360
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Patent No. 5721118
GENERAL INFORMATION:
                                                                                                                                                                                                                    1 AAAGTGCTGGGATTACAGG 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.87
Best Local Similarity 100.0
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                linear
MOLECULE TYPE: DN HYPOTHETICAL: NO US-08-629-939-10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: DN
HYPOTHETICAL: NO
US-08-759-873-10
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100.0%; Pred. No. 3.8e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: San Diego
STATE: California
COUNTRY: California
COUNTRY: United States
ZIP: 92122
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: BAPC compatible
COMPUTER: PatentIn Release #1.0, Version #1.25
GURRENT APPLICATION DATA:
RELING DATE:
FILING DATE:
                                                                                                                                                                                                               SOFTWARE: Patentin 1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/741,406
                  B: Campbell & Flores LLP
4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                          FILING DATE:
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/550,717
FILING DATE: 31-0CT-1995
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-UD 2317
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/741,406
FILING DATE:
APPLICATION NUMBER: US 08/550,717
FILING DATE: 31-OCT-1995
                                                                                                                                    COMPUTER READABLE FORM:
WEDIUM TYPE: Floppy disk
CMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2265 GTAGAGACAGGGTTTCACC 2283
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20 GTAGAGACAGGGTTTCACC 2
                                   STREET: 4370 La Jolla VI
CITY: San Diego
STATE: California
COUNTX: United States
ZIP: 92122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (619) 515-8949
INFORMATION FOR SEQ ID NO: 8
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.8%
Best Local Similarity 100.0
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
CORRESPONDENCE ADDRESS:
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US-08-741-406-8
                       ADDRESSEE:
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US-09-540-699-25/C
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OTHER INFORMATION: that is complementary to a portion of the human
OTHER INFORMATION: that is complementary to a portion of the human
NAME/KEY: misc_feature
INFORMATION: (19)
OTHER INFORMATION: /note= "n represents 5-DABCYL-thymidine"
US-09-540-699-23
                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.8%; Score 19; DB 1; Length 20; Best Local Similarity 100.0%; Pred. No. 3.8e+02; Matches 19; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 23, Application US/09540699;
Patent No. 6383752;
GENERAL INFORMATION;
APPLICANT: Agrawal, Sudhir;
APPLICANT: Agrawal, Sudhir;
TITLE OF INVENTION: Pseudo-Cyclic Oligonucleobases;
FILE REFERRNCE: 99,128-B;
CURRENT FLIING DATE: 2000-03-31;
PRIOR APPLICATION NUMBER: US 60/127,138;
PRIOR APPLICATION NUMBER: US 60/127,138;
PRIOR APPLICATION NUMBER: US 60/174,642;
NUMBER OF SEQ ID NOS: 26;
SOFTWARE: PALENT NOS: 26;
SOFTWARE: PALENT NOS: 26;
SEQ ID NO S: 26;
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-540-699-24/c
Sequence 24, Application US/09540699
Sequence 24, Application US/09540699
Sequence 24, Application US/09540699
SENERAL UNCRMANION:
APPLICANT: Agrawal, Sudhir
TITLE OF INVENTION: Pseudo-Cyclic Oligonucleobases
FILE REFERENCE: 99,128-B
CURRENT APPLICATION NUMBER: US/09/540,699
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGIGSTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: P-UD 2317
TELECOMMUNICATION INFORMATION:
TELEFONE: (619) 535-9001
TELEFAX: (619) 535-9001
TELEFAX: (619) 535-9001
TELEFAX: C19) 535-901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           677 GAGTGAGACAGGTGTCACC 696
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                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-540-699-23/c
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                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Artificial Sequence
ORGANISM: Artificial Sequence
ORGANISM: Artificial Sequence
ORGANISM: Artificial Sequence: Oligonucleotide
OTHER INFORMATION: that is complementary to a portion of the human
OTHER INFORMATION: MDM2 mRNA
NAME/KEY: misc_feature
LOCATION: (18)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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US-09-540-699-24
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US-09-540-699-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.8%; Score 19; DB 1; Length 20; Best Local Similarity 95.0%; Pred. No. 3.8e+02; Matches 19; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 19; DB 1; Length 20; 95.0%; Pred. No. 3.8e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-54/09-54/09-52/09-52/09-54/09-59-52/09-54/09-59-52/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-59/09-
; CURRENT FILING DATE: 2000-03-31
; PRIOR APPLICATION NUMBER: US 60/127,138
; PRIOR PILING DATE: 1999-03-31
; PRIOR FILING DATE: 2000-01-05
; PRIOR FILING DATE: 2000-01-05
; NUMBER OF SEQ ID NOS: 26
; SEQ ID NO 24
; LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 473
US-09-540-699-26/c
, Sequence 26, Application US/09540699
, Patent No. 6383752
, GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  676 TGAGTGAGAACAGGTGTCAC 695
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      674 TGTGAGTGAGAACAGGTGTC 693
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  20 TGNGTGAGAACAGGTGTCAC 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence PEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 95.03
Matches 19; Conservative
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                             COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/599,252
FILING DATE: 09-FEB-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MURASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFRENCE/DOCKET NUMBER: 29,959
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 887-0763
                             TITLE OF INVENTION: METHOD TO DIAGNOSE HEREDITARY
TITLE OF INVENTION: HEMOCHROMATOSIS
NUMBER OF SEQUENCES: 124
NUMBER OF SEQUENCES: ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 2000 Pennsylvania Ave. N.W., Suite 5500
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ?: 2000 Pennsylvania Ave. N.W., Suite 5500 Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COUNTRY: USA
ZIP: 20006-1888
COMPUTER REDABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/599,252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   us-08-599-252-38/C
; Sequence 38, Application US/08599252
; Patent No. 5705343
; GENERAL INFORMATION:
APPLICANT: PEDER, JOHN N.
APPLICANT: RIMMEL, BRUERS,
APPLICANT: THOMAS, WINNERS,
APPLICANT: THOMAS, WINNERS,
APPLICANT: THOMAS, WINNERS,
APPLICANT: THOMAS, WINNERS,
APPLICANT: HOMAS, WINNERS,
APPLICANT: HOMAS, WINNERS,
APPLICANT: HOMAS, WINNERS,
MURBER OF SEQUENCES: 124
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 90.9%; Pred. ...
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                                                                                                                                                                                                                 COUNTRY: USA
ZIP: 2006-1888
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
              APPLICANT: WOLFF, ROGER K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 22 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Best Local Similarity 90.9
Matches 20, Conservative
                                                                                                                                                                     CITY: Washington STATE: DC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
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Batent No. 6440737

GENERAL INFORMATION:
APPLICANT: Lex M. COWSERT
APPLICANT: Lex M. COWSERT
APPLICANT: Lex M. COWSERT
APPLICANT: SUSAN M. Freier
TITLE OF INVENTION: ANTISENSE MODULATION OF CELLULAR APOPTOSIS SUSCEPTIBILITY GENE 1
FILE REFREENTE: 2000-11-01
NUMBER OF SEQ ID NOS: 86
SEQ ID NO 17
LENGTH: 20
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OTHER INFORMATION: that is complementary to a portion of the human
OTHER INFORMATION: MDM2 mRNA
NAME/KEY: misc_feature
LOCATION: (14)
. LOCATION: (14)
. OTHER INFORMATION: /note= "n represents 5-DABCYL-thymidine"
US-09-540-699-26
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 19; DB 1; Length 20; 100.0%; Pred. No. 3.8e+02; tive 0; Mismatches 0; Indels
APPLICANT: Agrawal, Sudhir
APPLICANT: Kandimalla, Ekambar R.
TITLE OF INVENTION: Peeudo-Cyclic Oligonucleobases
FILE REPERENCE: 99,128-B
CURRENT APPLICATION NUMBER: US/09/540,699
CURRENT FILING DATE: 2000-03-31
PRIOR APPLICATION NUMBER: US 60/127,138
PRIOR APPLICATION NUMBER: US 60/174,642
PRIOR FILING DATE: 1999-03-31
PRIOR PILING DATE: 2000-01-05
NUMBER OF SEQ ID NOS: 26
SOFTWARE: Patentin Ver. 2.0
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               , OTHER INFORMATION: Antisense Oligonucleotide US-09-705-299-17
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Patent No. 5705343
GENERAL INFORMATION:
APPLICANT: PEDER, JOHN N.
APPLICANT: REBER, ANDREAS,
APPLICANT: KIMMEL, BRUCE E.
APPLICANT: THOMAS, WINSTON J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             672 TCTGTGAGTGAGAACAGGTG 691
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                                                                                                                                                                                                                                                                                                                                      TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Artificial Sequence
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Best Local Similarity 100.º
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-599-252-35/c
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US-08-874-186-11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.8%; Score 18.8; DB 1; Length 22; Best Local Similarity 90.9%; Pred. No. 3.8e+02; Matches 20; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-08-117-952-257/C

Sequence 257, Application US/08117952

Sequence 257, Application US/08117952

Sequence 257, Application US/08117952

SENERAL INFORMATION:

APPLICANT: Evans, Glen A.

APPLICANT: Smith, Michael W.

TITLE OF INVENTION: METHOD FOR GENERATION OF SEQUENCE

TITLE OF INVENTION: SAMPLED MAPS OF COMPLEX GENOMES

NUMBER OF SEQUENCES: 797

CORRESPONDENCE ADDRESS:

STREET: 444 SOUTH Flower Street, Suite 2000

STRIE CA

STRIE: CA
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MEDIUM TYPE: Floppy disk
COMPUTER: Eloppy disk
COMPUTER: El POPPy disk
COMPUTER: El PR PC COMPATIBLE
CORPERATION SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/117,952
FILING DATE: 07-SEP-1993
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/078,471
FILING DATE: 15-70N-1993
ATTORNEY/AGENT INFORMATION:
NAME: Reiter, Stephen E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Reiter, Stephen E.
REGISTRATION NUMBER: 31,192
REFERENCE/DOCKET NUMBER: P41 9423
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619-546-4737
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2113 GCTCTGTTACCCAGGCTGGAGT 2134
FILING DATE: 09-FEB-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: MUTAABLIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 9053-
TELECOMMUNICATION INFORMATION:
FELEX: 90-4030
INFORMATION FOR SEQ ID NO: 38:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
FYPE: nucleic acid
STRANDEDRESS: single
TYPE: TOOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     22 GCTCTATTGCCCAGGCTGGAGT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: Oligonucleotide
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 257: SEQUENCE CHARACTERISTICS: LENGTH: 22 base pairs TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 HYPOTHETICAL:
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US-08-117-952-257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-599-252-38
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Sequence 11, Application US/08874186
Fatent No. 5989885
GENERAL INFORMATION:
FAPLICANT: Taviglan, Sean V.
APPLICANT: Taviglan, Sean V.
APPLICANT: Taviglan, Sean V.
APPLICANT: Perry III, William L.
APPLICANT: SECINIC, Mark H.
TITLE OF INVENTION: SPECIFIC WITATIONS OF MAP KINASE KINASE
TITLE OF INVENTION: SUPPRESSOR IN VARIOUS TYPES OF CANCER
NUMBER OF SEQUENCES: 96
CORRESPONDENCE ADDRESS:
ADDRESSEE: Venable, Baetjer, Howard & Civiletti, LLP
STREET: 1201 New York Avenue, N.W., Suite 1000
CITY: Washington
STREET: D.
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Score 18.8; DB 1; Length 22;
Pred. No. 3.8e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Felease #1.0, Version #1.30 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/874,186 FILING DATE: CLASSIFICATION: 435 CLASSIFICATION: 435 APPLICATION NUMBER: US 08/782,482 FILING DATE: US 08/782,482 FILING DATE: 10-JAN-1997 ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NAME: Saxe, Stephen A.
REGISTRATION NUMBER: 38,609
REFERENCE/DOCKET NUMBER: 24884-121392-01
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-962-4848
TELEFAX: 202-962-8300
INFORMATION FOR SEQ ID NO: 11:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "Primer for STS."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: U.S.A.
ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                             2094 TTTTTTGAGACCGAGTCTTGCT 2115
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2261 TTTAGTAGAGACAGGGTTTCAC 2282
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                                                                                                                                        22 Tritrcsagacagacicriscr 1
    0.8%;
Query Match
Best Local Similarity 90.94
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Best Local Similarity 90.9
Matches 20; Conservative
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2; Indels
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CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
STREET: 2000 Pennsylvania Ave. N.W., Suite 5500
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OGER K.
METHOD TO DIAGNOSE HEREDITARY
HEMOCHROMATOSIS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 20006-1888
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
APPLICATION NUMBER: PCT/US96/06583
  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/06352
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match

0.8%; Score 18.8; DB 1;
Best Local Similarity 90.9%; Pred. No. 3.8e+02;
Matches 20; Conservative 0; Mismatches 2;
                                                                                                                    9053-0001.21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION UNDRAE:
APPLICATION UNDRAE:
ILING DATE: 09-FEB-1996
ATTORNEY/AGENT INFORMATION:
NAME: MURASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 9053-0001.2
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GGAGT 2134
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; Sequence 35, Application PC/TUS9606583
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22 gererafrececasser 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: BRAYNA, DENNIS T.
APPLICANT: FEDER, JOHN N.
APPLICANT: GNIRKE, ANDREAS
APPLICANT: THOMAS, WINSTON J.
APPLICANT: THOMAS, WINSTON J.
ITILE OF INVENTION: METHOD TO D
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2113 GCTCTGTTACCCAGGCT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; TOPOLOGY:
PCT-US96-06352-38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STATE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: DRAYNA, DENNIS T.
APPLICANT: FEDER, JOHN N.
APPLICANT: FEDER, JOHN N.
APPLICANT: GINEKE, ANDREAS
APPLICANT: KIMMEL, BRUCE E.
APPLICANT: THOMAS, WINSTON J.
APPLICANT: WOLFF, ROGER K.
TITLE OF INVENTION: HEMOCHROMATOSIS
NUMBER OF SEQUENCES: 124
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORRISON & FOERSTER
                                  APPLICANT: KIMMEL, BRUCE E.
APPLICANT: THOMAS, WINSTON J.
APPLICANT: WOLFF, ROGER K.
TITLE OF INVENTION: METHOD TO DIAGNOSE HEREDITARY
TITLE OF INVENTION: HEMOCHROMATOSIS
                                                                                                                                                                         ADDRESSEE: MORRISON & FOERSTER
STREET: 2000 Pennsylvania Ave. N.W., Suite 5500
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2000 Pennsylvania Ave. N.W., Suite 5500
                                                                                                                                                                                                                                                    COUNTRY: USA
ZIP: 20006-1888
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRITA APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/06352
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   9053-0001.21
                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 09-FEB-1996
ATTOKNEY/AGENT INFORMATION:
NAME: WURASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 9053-0001.2
TELEFAN: (202) 897-1500
TELEFAN: (202) 897-0763
TELEFAX: (202) 897-0763
TELEFAX: (202) 897-0763
TELEFAX: 90-4030
INFORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDENNES: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22 GCTCTATTGCCCAGGCTGGAGT 1
JOHN N.
, ANDREAS
                                                                                                                                          124
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Best Local Similarity 90.9°
Matches 20; Conservative
                                                                                                                                    NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 20006-1888
COMPUTER READABLE FORM
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CITY: Washington
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: DC COUNTRY: U
                                                                                                                                                                                                                  CITY: Wash
STATE: DC
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PCT-US96-06352-35
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Gaps

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Length 22;

22 GCTCTATTGCCCAGGCTGGAGT

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(202) 887-1500

TELEPHONE:

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MOLECULE TYPE:
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US-08-222-177A-341
                                                                                          US-08-031-143B-25
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                                                                                                                                                                                                                 Length 22;
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                                                                                                                                                                                                                                                         2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: DRAYNA, DENNIS T.
APPLICANT: FREDER, JOHN N.
APPLICANT: GNIRKE, ANDREAS
APPLICANT: THOMAS, WINSTON J.
APPLICANT: THOMAS, WINSTON J.
APPLICANT: WOLFF, ROGER K.
TITLE OF INVENTION: HEMOCHROMATOSIS
NUMBER OF SEQUENCES: 124
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: MORRISON & FOERSTER
STREET: 2000 Pennsylvania Ave. N.W., Suite 5500
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US96/06583
                                                                                                                                                                                                            Query Match
0.8%; Score 18.8; DB 1;
Best Local Similarity 90.9%; Pred. No. 3.8e+02;
Matches 20; Conservative 0; Mismatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/599,252
FILING DATE: 09-FEB-1996
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                               2113 GCTCTGTTACCCAGGCTGGAGT 2134
                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 38, Application PC/TUS9606583 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: USA
ZIP: 20006-1888
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                     22 GCTCTATTGCCCAGGCTGGAGT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: MUTASHIGE, KATE H.
REGISTRATION NUMBER: 29,959
REFERENCE/DOCKET NUMBER: 9053
TELEPONE: (202) 887-053
TELEPONE: (202) 887-053
TELER: 90-4030
INFORMATION FOR SEQ ID NO: 38:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDENNESS: single
TELERA: (202) 887-0763
TELEX: 90-4030
INFORMATION FOR SEQ ID NO: 35
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 90.99
Matches 20; Conservative
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                                                                                                                                                   linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:
                                                                                                                                                   ; TOPOLOGY:
PCT-US96-06583-35
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TOPOLOGY:
PCT-US96-06583-38
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2113 GCTCTGTTACCCAGGCTGGAGT 2134

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Sequence 341, Application US/08222177A

Patent No. 5582379

GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: (dc-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
TITLE OF SEQUENCES:
NUMBER OF SEQUENCES:
OCRRESPONDENCE ADDRESS:
ADDRESSEE: DeWitt Ross & Stevens, S.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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Sequence 25, Application US/08031143B
| Patent No. 5518880
| GENERAL INFORMATION: APPLICANT: MCBRIDS: O. WESLEY
| TITLE OF INVENTION: METHODS FOR DIAGNOSIS AND | TITLE OF INVENTION: METHODS FOR SCIENCES: ORRESPONDENCE ADDRESS: ADDRESSEE: MORGAN & FINNEGAN | STREET: 345 PARK AVE. | CULTY: NEW YORK | STATE: NEW YORK | COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SOUTWARE: WORD PERFECT # 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/031,143B
FILING DATE: 12-MAR-1993
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: WILLIAM S. FEILER
REGISTRATION NUMBER: 26,728
REFERENCE/DOCKET NUMBER: 26,728
TELECHONISICATION INFORMATION:
TELECHONE: 212-758-4800
TELECHONE: 212-758-4800
                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2353 ATTACAGGCATGAGCCACCG 2372
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    25:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 421792
INPORMATION FOR ESQ ID NO: SEQUENCE CHARCTERISTICS: LENGTH: 20
TYPE: NUCLEIC ACID STRANDEDNESS: SINGLE TOPOLOGY: UNKNOWN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 95.0
Matches 19; Conservative
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REFERENCE/DOCKET NUMBER: 07265/079001
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             La Jolla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: 1
STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; Sequence 71, 7, Application US/08588821
; Patent No. 5712097
; GENERAL INFORMATION:
APPLICANT: Kern, Scott E.
TITLE OF INVENTION: NOVEL TUMOR SUPPRESSOR GENE, DPC4
NUMBER OF SEQUENCES: 91
; CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
; STREET: 4225 Executive Square, Suite 1400
; CITY: La Jolla
STATE: CA COUNTRY: USA
; COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATURG SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/588,821
FILING DATE: 19-UAN-1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Haile, Lise A.
REGISTRATION NUMBER: 38,347
                                                                              ZIP: 53717-1914

CMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                         FILING DATE:
CLASSIFICATION: 435
PRIOR APPLICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION: G08) 831-2100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2339 CCCAAAGTGCTGGGATTACA 2358
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     INFORMATION FOR SEQ ID NO: 341:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                  Madison
Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; IMMEDIATE SOURCE;
; CLONE: mfd107p1
US-08-222-177A-341
                                                              USA
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                                                              COUNTRY:
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                                                                                                                                                                                                                                                                                                                                        Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 70, Application US/08915214
; Sequence 70, Application US/08915214
; Patter No. 5814457
; GENERAL INFORMATION:
APPLICANT: Hahn, Stephan A.
ITILE OF INVENTION: NOVEL TUMOR SUPPRESSOR GENE, DPC4
; UNMBER OF SEQUENCES: 91
; CORRESPONDENCE ADDRESS:
ADDRESSEE Fish & Richardson P.C.
STREET: 4225 Executive Square, Suite 1400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: LA

COUNTRY: USA

ZIP: 92037

CONFUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
CONFUTER: IBM PC compatible
CONFUTER: IBM PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/915,214
FILING DATE: 20-AUG-1997
CLASSIFICATION 1435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/588,821
FILING DATE: 19-JAN-1996
ATTORNEY/AGENT INFORMATION:
NAME: Haile, Lisa A.
REGISTRATION NUMBER: 07265/079001
TELECOMMUNICATION INFORMATION:
TELEPAN: 619/678-5070
TELECOMMUNICATION INFORMATION:
TELEPAN: 619/678-5030
TELECOMMUNICATION INFORMATION:
TELEPAN: 619/678-5030
TELECOMMUNICATION INFORMATION:
TELEPAN: 619/678-5030
TELECOMMUNICATION INFORMATION:
TELEPAN: 619/678-5030
TELEPAN: 619/678-5030
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                  0.8%; Score 18.4; DB 1;
95.0%; Pred. No. 4.1e+02;
tive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                              2338 TCCCAAAGTGCTGGGATTAC 2357
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TELECOMMUNICATION INFORMATION:
TELEPHONE: 619/678-5070
TELEPHONE: 619/678-5099
INFORMATION FOR SEQ ID NO: 70:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
US-08-588-821-70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 rcccaaagrecreegarrrc 20
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Best Local Similarity 95.07
Matches 19; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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COMPUTER READABLE FORM:
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Linear
No
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ANTI-SENSE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: EXPRESSION
INTERED PINVENTION: EXPRESSION
INTERED PROPERS: 271
CORRESPONDENCE ADDRESS:
ADDRESSE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
COMPUTER READABLE FORM:
US-09-005-532-70

Sequence 70, Application US/09005532

Patent No. 595229

GENERAL INFORMATION:

APPLICANT: Hahn, Stephan A.

ITLE OF INVENTION: NOVEL TUMOR SUPPRESSOR GENE, DPC4

NUMBER OF SEQUENCES: 91

CORRESPONDENCE ADDRESS:

ADDRESSE: Fish & Richardson P.C.

STARE: 4225 Executive Square, Suite 1400

CITY: La Jolla

STATE: CA.
                                                                                                                                                                                                                                                                                                   ZOUNTIN USA
ZIP: 92037
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/005,532
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILLING JAPE

CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/588,821
FILLING DATE: 19-JAM-1996
ATTORNEY/AGENT INFORMATION:
NAME: Halle, Lisa A.
REGISTRATION NUMBER: 07265/079001
TELEPHONE: 619/678-5070
TELEPHONE: 619/678-5099
INFORMATION FOR SEQ ID NO: 70:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDBRSS: single.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2338 TCCCAAAGTGCTGGGATTAC 2357
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; Sequence 27, Application US/09280805
; Patent No. 6184212
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US-09-005-532-70
                                                                                                                                                                                                                                                                                        USA
                                                                                                                                                                                                                                                                                      COUNTRY:
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Sequence 12, Application US/08430225A

Sequence 12, Application US/08430225A

Patent No. 6204000

GENERAL INFORMATION:
APPLICANT: Dong, Jin-Tang; Barrett,
APPLICANT: Dong, Jin-Tang; Barrett,
APPLICANT: J. Carl; Lamb, Patricia W.; Isaacs, John T.
TITLE OF INVENTION: DIAGNOSTIC METHODS AND
TITLE OF INVENTION: HUMAN METASTASIS SUPPRESSOR GENE KAII
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: MORGAN & FINNEGAN, L.L.P.
STREET: 345 PARK AVENUE
CITY: NEW YORK
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ..
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MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE COMPUTER: IBM PC OPERATING SYSTEM: WINDOWS 95
                                              CORFOLER: LEGY WINDOWS 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 09/048,810
FILING DATE: March 26, 1998
ATTONNEY AGENT INFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0346
TELECHONE: 609-810-1515
TELECHONE: 609-810-1515
TELECHONE: CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WORDPERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/430,225A
FILING DATE: 28-APR-195
CLASSIFICATION: 514
ATTORNEY/AGENT INFORMATION:
NAME: RICHARD W. BORK
REGISTRATION NUMBER: 36,459
REFERENCE/DOCKET NUMBER: 2026-4172
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECHARD: (212) 751-6849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1695 TTTACATGTGCAAAGAAGCT 1714
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OTHER INFORMATION: Antisense Oligonucleotide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 Arcrerakerrerakie 20
                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 492
US-09-733-294A-82
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                                                                                                                   Score 18.4; DB 1; Length 20;
Pred. No. 4.1e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0.8%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 4.1e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                             Sequence 27, Application US/09048810
Patent No. 6238921
GENERAL INFORMATION:
APPLICANT: Graham, Brett P. Monia
TITLE OF INVENTION: ANTISENSE OLIGONUCLEOTIDE
TITLE OF INVENTION: MODILATION OF HUMAN MDM2 EXPRESSION
NUMBER OF SEQUENCES: 32
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 08053

COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM 486
OPERATING SYSTEM: WINDOWS FOR WORKGROUPS
SOFTWARE: WORDPERFET 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/048,810
FILING DATE: herewith
CLASSIFICATION: ATTORNEY FORM TINFORMATION:
NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0302
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                 2179 TTCGCACCATTCTCCTGCCT 2198
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1695 TTTACATGTGCAAAGAAGCT 1714
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                                                                                                                                                                                                                           1 TTCACACCATTCTCCTGCCT 20
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                                                                                                                     0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELEFAX: 609-810-1454
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                               Query Match
Best Local Similarity 95.09
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.8
Best Local Similarity 95.0
Matches 19; Conservative
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: Linear
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Marlton
STATE: NJ
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                 US-09-048-810-27/c
                                                          ; TOPOLOGY:
US-08-430-225A-12
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TITLE OF INVENTION: Antisense Fucosyltransferase Sequences and Methods of TITLE OF INVENTION: Use Thereof FILE REFERENCE: Weston and Hiller CURRENT APPLICATION NUMBER: US/09/556,031
CURRENT FILING DATE: 2000-04-20
PRIOR PILING DATE: 1999-04-26
NUMBER OF SEQ ID NOS: 24
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 18
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                             ) OTHER INFORMATION: Description of Artificial Sequence:antisense; OTHER INFORMATION: Oligonucleotide
US-09-556-031-18
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GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: William Gaarde
APPLICANT: William Gaarde
APPLICANT: William Garde
TITLE OF INVENTION: ANTISENSE MODULATION OF TERT EXPRESSION
FILE REFERENCE: ISPH-0527
CURRENT APPLICATION NUMBER: US/09/733,294A
CURRENT FILING DATE: 2000-12-07
PRIOR APPLICATION NUMBER: 09/572,423
FRIOR APPLICATION NUMBER: 09/572,423
NUMBER OF SEQ ID NOS: 108
SEQ ID NO 82
LENGTH: 20
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Fatent No. 6743909
GENERAL INFORMATION:
APPLICANT: Lex M. Cowsert
APPLICANT: Kenneth W. Dobie
TITLE OF INVENTION: ANTISENSE MODULATION OF PTPN12 EXPRESSION
FILE REPERENCE: PTS-0016
CURRENT FILING DATE: 2002-06-17
NUMBER OF SEQ ID NOS: 123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.8%; Score 18.4; DB 1; Length 20; 95.0%; Pred. No. 4.1e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match

0.8%; Score 18.4; DB 1;
Best Local Similarity 95.0%; Pred. No. 4.1e+02;
Matches 19; Conservative 0; Mismatches 1;
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APPLICANT: THE GOVERNMENT OF THE UNITED STATES OF AMERICA AS
APPLICANT: REPRESENTED BY THE SECRETARY, DEPARTMENT OF HEALTH AND HUMAN
APPLICANT: SERVICES
APPLICANT: STATUS OF TECHNOLOGY TRANSFER, NATIONAL
APPLICANT: INSTITUTES OF HEALTH, BOX OTT, BETHESDA, MARYLAND 20892 USA
TITLE OF INVENTION: METHODS FOR DIAGNOSIS AND TREATMENT OF
TITLE OF INVENTION: XSCID
NUMBER OF SEQUENCES: 69
ONRESPONDENCE ADDRESS:
ADDRESSE: ADDRESS:
ADDRESSE: ADDRESS :
STREET: 345 PARK AVE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 1, Application US/09104497
Sequence 1, Application US/09104497
Batent No. 6028245
GENERAL INFORMATION:
APPLICANT: WASYLYK, Bohdan
APPLICANT: TOCQUE, Bruno
APPLICANT: ALKHALAF, Moussa
TITLE OF INVENTION: TRANSGENIC ANIMALS OVEREXPRESSING MDM2
FILE REFERENCE: A2716A-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 18.4; DB 1;
Pred. No. 4.1e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: WORD PERFECT # 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/02891
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2026-4061
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/031,143
FILING DATE: 12-MAR-1993
APPLICATION NUMBER: 08/121,435
FILING DATE: 14-SEPT-1993
ATTORNEY/AGENT INFORMATION:
NAME: WILLIAM S. FEILER
REGISTRATION NUMBER: 26,728
REFERENCE/DOCKET NUMBER: 2026-406
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 12-758-4800
Application PC/TUS9402891
                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDIUM TYPE: FLOPPY DISK
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: WORD PERPECT # 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2353 ATTACAGGCATGAGCCACCG 2372
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LET: 69
LEET: ADDRESS:
LEET: 345 PARK AVE.
': NEW YORK
'Y: USA
'15'
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DESCRIPTION: OLIGONUCLEOTIDE POTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     25:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ORGANISM: HUMAN
INDIVIDUAL ISOLATE: IL-2R
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0.8%;
ilarity 95.0%;
Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 212-751-6849
TELEX: 421792
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH: 20
TYPE: NUCLEIC ACID
STRANDEDNESS: SINGLE
TOPOLOGY: UNKNOWN
                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 10154
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 19; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              HYPOTHETICAL: 1
ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORIGINAL SOURCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PCT-US94-02891-25
                                                                                                                                                                                                                                                                                                                                            STATE: NE COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 JS-09-104-497-1
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J. Carl; Lamb, Patricia W.; Isaacs, John T.
JITLE OF INVENTION: DIAGNOSTIC METHODS AND
GENE THERAPY USING REAGENTS DERIVED FROM THE
HUMAN METASTASIS SUPPRESSOR GENE KALI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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                                                                                                                                                                     Query Match 0.8%; Score 18.4; DB 1; Length 20; Best Local Similarity 95.0%; Pred. No. 4.1e+02; Matches 19; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match 0.8%; Score 18.4; DB 1; Length 20; Best Local Similarity 95.0%; Pred. No. 4.1e+02; Matches 19; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: USA

ZIP: 10154

COMPUTER READABLE FORM:
MEDIUM TYPE: PLOPPY DISK

MEDIUM TYPE: PLOPPY DISK

COMPUTER: ISM PC COMPATTBLE

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: MICROSOFT WORD 97

CURENT APPLICATION DATA:
APPLICATION NUMBER: US/09/795,380

FILING DATE: 27-Feb-2001

PRIOR APPLICATION NUMBER: 09/232,507

FILING DATE: <Unknown>
ATTORNEY/AGBRT INFORMATION:
NAME: RICHARD W. BORK

REGISTRATION NUMBER: 36,459

REFERENCE/DOCKET NUMBER: 36,459

TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFFAKE (212) 751-6849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: MORGAN & FINNEGAN, L.L.P. STREET: 345 PARK AVENUE CITY: NEW YORK STATE: NEW YORK COUNTRY: USA
                                                                                               ; OTHER INFORMATION: Antisense Oligonucleotide US-10-172-911-80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 12:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2179 TICGCACCATICICCIGCCI 2198
                                                                                                                                                                                                                                                                    2120 TACCCAGGCTGGAGTGCAGT 2139
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 12, Application US/09795380 Patent No. 6756201 GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                           TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 495
PCT-US94-02891-25
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     LENGTH: 20
                                                                             FEATURE:
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Gaps

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Indels

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Length 20;

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US-09-784-423-80/C
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Sequence 80, Application US/09018584A

Patent No. 628863

GENERAL INFORMATION:

APPLICANT: Schumm, James W.

TITLE OF INVENTION: MATERALS AND METHODS FOR

TITLE OF INVENTION: IDENTIFYING AND ANALYZING INTERMEDIATE TANDEM

TITLE OF INVENTION: REPEAT DNA MARKERS

TITLE OF SEQUENCES: 147
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                                                                                                                                                                                                                                       0.8%; Score 18.4; DB 1; Length 23; 95.0%; Pred. No. 3.9e+02; tive 0; Mismatches 1; Indels
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ZIP: 53711-5399
COMPUTER READABLE FORM:
MEDIUM TYBE: Diskette - 3.5 inch, 1.44 Mb
COMPUTER: IBM compatible PC
OPERATING SYSTEM: Windows 95
SOFTWARE: Word 97 (DOS text format)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/018,584A
FILING DATE: U4-Feb-1998
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Grady J. Frenchick
REGISTRATION NUMBER: 29,018
REFERENCE/DOCKET NUMBER: 16026.9180
TELECOMMUNICATION:
CURRENT APPLICATION NUMBER: US/09/104,497
CURRENT FILING DATE: 1998-06-25
EARLIER APPLICATION NUMBER: 60/051,739
EARLIER FILING DATE: 1997-07-03
NUMBER OF SEQ ID NOS: 7
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 1
LENGTH: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 2800 Woods Hollow Road CITY: Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2187 ATTCTCCTGCCTCAGCCTCC 2206
                                                                                                                                                                                                                                                                                                                      312 ATGTGCAATACCAACATGTC 331
                                                                                                                                                                                                                                                                                                                                                 4 ATGGGCAATACCAACATGTC 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (608) 257-2275
INFORMATION FOR SEQ ID NO: 8
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                       Query Match
Best Local Similarity 95.0°
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 95.0
Matches 19; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SS: Single
Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Promega C
                                                                                                                                                           TYPE: DNA ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: Nucleic Acid STRANDEDNESS: Sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TOPOLOGY:
US-09-018-584A-80
                                                                                                                                                                                                  US-09-104-497-1
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RESULT 498

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                                                                APPLICANT: Schumm, James W.

Bacher, Jeffery W.

TITLE OF INVENTION: MATERIES AND METHODS FOR
REPEAT DNA MARKERS

REPEAT DNA MARKERS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            1; Indels
                                                                                                                                                                                                                                                                                          STATE: Wisconsin
STATE: Wisconsin
COUNTRY: U.S.A.
21P: 53711-539
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb
COMPUTER: IBM comparable PC
OOMPUTER: IBM comparable PC
OOPERATING SYSTEM: Windows 95
SOFTWARE: Word 97 (DOS text format)
APPLICATION DATA
APPLICATION NUMBER: US/09/784,423
FILING DATE: 15-Feb-2001
CLASSIFICATION: cUnknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            In-Situ Hybridization Probes for
Identification and Banding of
Specific Human Chromosomes and
Regions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 18.4; DB 1;
Pred. No. 3.9e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:

APPLICATION NUMBER: 09/018,584
FILING DATE: 04-Feb-1998
ATTORNEY/AGENT INFORMATION:
NAME: Grady J. Frenchick
REGISTRATION NUMBER: 29,018
RERERENCE/DOCKET NUMBER: 16026.9180
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 257-2275
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
0.8%; Score 18.4; D
Best Local Similarity 95.0%; Pred. No. 3.9e
Matches 19; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: Linear
SEQUENCE DESCRIPTION: SEQ ID NO: 80
                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSER: Promega Corporation
STREET: 2800 Woods Hollow Road
CITY: Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     E: Arnold, White & Durkee P.O. Box 4433
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-070-517-1

Sequence 1, Application US/08070517

Sequence 1, Application US/08070517

Parent No. 553869

APPLICANT: Michael J. Siciliano
APPLICANT: Pu Liu
TITLE OF INVENTION: In-Situ Hybrid
TITLE OF INVENTION: Identification
TITLE OF INVENTION: Regions
TITLE OF INVENTION: Regions
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:

ADDRESSEE: Arnold, White & Durke
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2187 ATTCTCCTGCCTCAGCCTCC 2206
Sequence 80, Application US/09784423
Patent No. 6767703
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   22 Arricriciriscircascrici
                                                                                                                                                                                   NUMBER OF SEQUENCES: 147
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: (608) 257...
INFORMATION FOR SEQ ID NO: 80
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CITY: Houston
STATE: Texas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Texas
: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-784-423-80
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STREET:
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NAME: Granahan, Patricia
REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: MIT-
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-861-6240
TELEFAX: 617-861-9540
                DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: Two Militia Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Lexington
STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
            MOLECULE TYPE:
HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY: US
                                                                                         US-08-118-441-1
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APPLICANT: Gilliam, T. Conrad
TITLE OF INVENTION: GENE
NUMBER OF SEQUENCES: 29
CORRESPONDENCES: 29
CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooper & Dunham
STREET: 30 ROCKéfeller Plaza
CITY: New York
STATE: New York
COUNTRY: United States of America
ZIP: 1012
ZIP: 1012
COMPUTER: Floppy disk
COMPUTER: Floppy disk
COMPUTER: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/118,441
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CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION UNDER: 28,678
REFERENCE/DOCKET NUMBER: 0555/44011
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 977-9550
TELEPHONE: (212) 664-0525
TELER: 42253 COOP UI
SEQUENCE CHARACTERISTICS:
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 UTSC:290/KIT
                                                                                                                                          CURRENT APPLICATION DATA;

APPLICATION NUMBER: US/08/00.517

FILING DATE: 19930601

CLASSIFICATION NUMBER: US/08/070,517

FILING DATE: 19930601

CLASSIFICATION: 435

ATTORNEY/AGENT INFORMATION:
NAME: Barbara S. Kitchell:
REGISTRATION NUMBER: UTSC:290/KI'
FELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 320-7200
TELEPHONE: (512) 374-7577

INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
LEGENCE CHARACTERIST
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy Disk
CMEDUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2351 GGATTACAGGCATGAGCCA 2369
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Patent No. 5578493
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 GGATTACAGGYRTGAGCCA 19
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STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-070-517-1
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Gaps
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Query Match 0.8%; Score 18.2; DB 1; Length 19; Best Local Similarity 89.5%; Pred. No. 4.2e+02; Matches 17; Conservative 2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Brook, J. David
APPLICANT: Brook, J. David E.
APPLICANT: Shaw, Duncan J.
APPLICANT: Shaw, Duncan J.
APPLICANT: Johnson, Keith J.
TITLE OF INVENTION: DNA SEQUENCE ENCODING THE MYOTONIC
TITLE OF INVENTION: DYSTROPHY GENE AND USES THEREOF
TORKESPONDENCES: 14
CORRESPONDENCES: 14
CORRESPONDENCES: Hamilton, Brook, Smith & Reynolds, P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ZIP: 02713
ZIP: 02713
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
APPLICATION NUMBER: US/08/422,699A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRICATION DATA:
PRICATION APPLICATION DATA:
APPLICATION NUMBER: US 08/023,612
FILING DATE: 26-FEB-1993
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US 07/839,255
FILING DATE: 20-FEB-1992
PRICA APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/01545
FILING DATE: 19-FEB-1993
PRICA APPLICATION DATA:
APPLICATION NUMBER: PCT/GB93/00253
FILING DATE: 05-FEB-1993
PRICA APPLICATION NUMBER: BCT/GB93/00253
FILING DATE: 06-FEB-1993
APPLICATION NUMBER: GB9202485.0
FILING DATE: 06-FEB-1993
APPLICATION NUMBER: GB9202485.0
FILING DATE: 06-FEB-1993
APPLICATION NUMBER: GB9202485.0
FILING DATE: 06-FEB-1993
                                                                                                                                                                                                                                                                                             US-08-422-699A-13
; Sequence 13, Application US/08422699A
Patent No. 5955.65
; GENERAL INFORMATION:
                                                                                                                    2351 GGATTACAGGCATGAGCCA 2369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION BATA:
APPLICATION NUMBER: 08/422,706
FILING DATE:
                                                                                                                                                      1 GGATTACAGGYRTGAGCCA 19
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linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STRANDEDNESS:
                            US-08-422-706B-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-08-338-579A-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            RESULT 503
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                                                                                                                                                Query Match 0.8%; Score 18.2; DB 1; Length 19; Best Local Similarity 89.5%; Pred. No. 4.2e+02; Matches 17; Conservative 2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 13. Application US/08422706B
Patent No. 597733
GENERAL INFORMATION:
APPLICANT: Horok, J. David
APPLICANT: Harley, Helen G.
APPLICANT: Harley, Helen G.
APPLICANT: Harley, Helen G.
APPLICANT: Harley, Helen G.
APPLICANT: J. HELEN G.
APPLICANT: J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOCTAMER: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/422,706B
FILING DATE: 14-APR-1995
CLASSIFICATION DATA:
APPLICATION NUMBER: US 08/23,612
FILING DATE: 26-FEB-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/839,255
FILING DATE: 20-FEB-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/01545
FILING DATE: 20-FEB-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/GB93/0253
FILING DATE: 19-FEB-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: MT-5830A2
FILING DATE: 05-FEB-1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: MIT-5830A2
FILING DATE: 06-FEB-1993
FRIOR APPLICATION NUMBER: MIT-5830A2
FILING DATE: 06-FEB-1993
FRIOR APPLICATION NUMBER: MIT-5830A2
FILING BATE: 06-FEB-1993
FRIEBEPHONE: 617-861-6240
INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
FILINGTH: 19 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Hamilton, Brook, Smith & Reynolds, P.C. STREET: Two Militia Drive CITY: Lexington STATE: Massachusetts COUNTRY: US
                                                                                                                                                                                                                                                                                                                                  2351 GGATTACAGGCATGAGCCA 2369
                                                                                                                                                                                                                                                                                                                                                                                          1 GGATTACAGGYRTGAGCCA 19
, MOLECULE TYPE: DNA (genomic) US-08-422-699A-13
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                                                                                 Gaps
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0.8%; Score 18.2; DB 1; Length 19; 89.5%; Pred. No. 4.2e+02; tive 2; Mismatches 0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                Sequence 1, Application US/08338579A

Sequence 1, Application US/08338579A

Patent No. 6068975

GENERAL INFORMATION:
FAPLICANT: Tanzi, Rudolph E.

TITLE OF INVENTION: ISOLATION AND USES OF A WILSON'S

TITLE OF INVENTION: ISOLATION AND USES OF A WILSON'S

TITLE OF INVENTION: ISOLATION AND USES OF A WILSON'S

TITLE OF INVENTION: DISEASE GENE
NUMBER OF SEQUENCES: 107

CORRESPONDENCE ADDRESS:

ADDRESSEE: COOPER & Dunham
STREET: 188 Avenue of the Americas

CITY: New York
STATE: New York
STATE: New York

COUNTRY: United States of America'

COMPUTER: New York

STATE: New York

COMPUTER: New York

STATE: New York

COMPUTER: New York

STATE: New York

STATE: New York

COMPUTER: New York

STATE: New York
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Outlier 1, DB 1;
Best Local Similarity 89.5%; Pred. No. 4.2e+02;
Matches 17; Conservative 2; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2351 GGATTACAGGCATGAGCCA 2369
                                                                                                                                                       2351 GGATTACAGGCATGAGCCA 2369
                                                                                                                                                                                                  1 GGATTACAGGYRTGAGCCA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 GGATTACAGGYRTGAGCCA 19
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   Query Match 0.8
Best Local Similarity 89.5
Matches 17; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 19 base pairs
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                                                                                                                                                                                                                       Score 18.2; DB 1; Length 19;
Pred. No. 4.2e+02;
2; Mismatches 0; Indels
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Pred. No. 4.2e+02;
2; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                       PCT-US94-09851-1

Sequence 1, Application PC/TUS9409851

GENERAL INFORMATION:

APPLICANT: Gilliam, T. Conrad

APPLICANT: Tanzi, Rudolph E.

TITLE OF INVENTION: ISOLATION AND USES OF A WILSON'S

TITLE OF INVENTION: DISEASE GENE

NUMBER OF SEQUENCES: 92

CORRESPONDENCE ADDRESS:

ADDRESSEB: Cooper & Dunham
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSER: Cooper & Dunham
STREET: 30 Rockefeller Plaza
CITY: New York
STATE: New York
COUNTRY: United States of America
ZIP: 10112
COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/09851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE:

CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REGISTRATION NUMBER: 0575/44011-PCT
TELECOMMUNICATION INFORMATION:
TELEPAX: (212) 977-9550
TELEPAX: (212) 64-0525
TELEFAX: (212) 64-0525
TELEFAX: (212) 64-0525
TELERAY: (212) 64-0525
TELERAY: (212) 64-0525
TELEX: 422523 COOP UI
SEQUENCE CHARACTERISTICS:
FILE REFERENCE: Davies Col
CURRENT APPLICATION NUMBER: US/09/078,294
UCHRENT FILING DATE: 1998-05-13
NUMBER OF SEQ ID NOS: 29
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                        2351 GGATTACAGGCATGAGCCA 2369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2351 GGATTACAGGCATGAGCCA 2369
                                                                                                                                                                                                                                                                                                                            1 GGATTACAGGYRTGAGCCA 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GGATTACAGGYRTGAGCCA 19
                                                                                                                                                                                                                     0.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.8%;
Best Local Similarity 89.5%;
Matches 17; Conservative
                                                                                                                                                                                                                   Query Match
Best Local Similarity 89.5°
Matches 17, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 19 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                           TYPE: DNA
ORGANISM: DNA primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
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                                                                                                                                                                                  US-09-078-294-1
                                                                                                    SEQ ID NO 1
LENGTH: 19
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RESULT 506

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APPLICANT: Mackenzie, Alex E.
APPLICANT: Korneluk, Robert G.
APPLICANT: Mandevan, Mani S.
APPLICANT: McLean, Michael
APPLICANT: McLean, Michael
APPLICANT: NcLean, Michael
APPLICANT: Roy, Natalie
APPLICANT: Roy, Natalie
TITLE OF INVENTION: Neuronal Apoptosis Inhibitor Protein, Gene Sequence and
FILE REFERENCE: 3477-112, 033477/139914
FILE REFERENCE: 3477-112, 033477/139914
CURRENT APPLICATION NUMBER: US/08/836,134A
CURRENT APPLICATION NOS: 23
SOFTWARE: PatentING DATE: 1997-06-20
SEQ ID NO 16
LENGTH: 23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Mackenzie, Alex B.
APPLICANT: Mackenzie, Robert G.
APPLICANT: Mackenzie, Robert G.
APPLICANT: Mackenzie, Mani S.
APPLICANT: Machen, Michael
APPLICANT: Machen, Michael
APPLICANT: Ikeda, Joh-e
APPLICANT: Ikeda, Joh-e
APPLICANT: Neuronal Apoptosis Inhibitor Protein, Gene Sequence and
TITLE OF INVENTION: Nutations Causative of Spinal Muscular Atrophy
FILE REPERBNCE: 3477-112, 033477/139914
CURRENT APPLICATION NUMBER: US/09/493,784
CURRENT FILING DATE: 2000-01-28
FRIOR APPLICATION NUMBER: 08/836,134
FRIOR APPLICATION NUMBER: 08/836,134
FRIOR APPLICATION NUMBER: 08/836,134
SROF FILING DATE: 1997-06-20
NUMBER OF SEQ ID NOS: 23
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 16
LENGTH: 23
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Description of Artificial Sequence: PCR primer
US-08-836-134-16
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US-09-493-784-16
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2227 CATCTGCCACCACCTGGCTAA 2249
Sequence 16, Application US/08836134A Patent No. 6020127 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 16, Application US/09493784; Patent No. 6429011; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Artificial Sequence
PEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     US-09-493-784-16/c
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Nucleic Acid
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Sequence 96, Application US/09018584A
Sequence 96, Application US/09018584A
PREDICANT: Schumm, James W.
APPLICANT: Bacher, Jeffery W.
TITLE OF INVENTION: IDENTIFYING AND MALYZING INTERMEDIATE TANDEM
TITLE OF INVENTION: IDENTIFYING AND ANALYZING INTERMEDIATE TANDEM
TITLE OF INVENTION: REPEAT DNA MARKERS
NUMBER OF SEQUENCES: 147
CORRESPONDENCE ADDRESS:
ADDRESSEE: Promega Corporation
STREET: 2800 Woods Hollow Road
CITY: Madison
STREET: 2800 Woods Hollow Road
CITY: Madison
COMPUTE: Diskette - 3.5 inch, 1.44 Mb
COMPUTE: IBM Compatible PC
OPERATING SYSTEM: Windows 95
SOUTHWARE: US MONT 97 (DOS text format)
CURRENT APPLICATION NUMBER: US/09/018,584A
FILIKO DATE:
CHARLE DATES TO A FEB-1998
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                                Sequence 167. Application US/09544398B

Patent No. 6770461
GENERAL INFORMATION:
APPLICANT: Carulli, John P.
APPLICANT: Little, Randall D.
APPLICANT: Little, Robert R.
APPLICANT: Johnson, Mark L.
TITLE OF INVENTION: High bone mass gene of 11q13.3
FILE REPERBANCE: 032796-013
CURRENT APPLICATION NUMBER: US/09/544,398B
                                                                                                                                                                                                                                                                            PRIOR APPLICATION NUMBER: US 09/229,319
PRIOR FILING DATE: 1999-01-13
PRIOR FILING DATE: 1998-01-13
PRIOR FILING DATE: 1998-01-13
PRIOR PILING DATE: 1998-01-13
PRIOR PILING DATE: 1998-10-23
NUMBER OF SEQ ID NOS: 641
SEQ ID NO 167
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Grady J. Frenchick
REGISTATION NUMBER: 29,018
REFERENCE/DOCKET NUMBER: 16026.9180
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2259 CTTTTAGTAGAGACAGGGTTTCA 2281
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TELEFAX: (608) 257-2275
INFORMATION FOR SEQ ID NO: 96
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.8%
Best Local Similarity 87.0%
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-544-398B-167
                  -09-544-398B-167
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                                                                                      Query Match 0.8%; Score 18.2; DB 1; Length 24; Best Local Similarity 87.0%; Pred. No. 4e+02; Matches 20; Conservative 0; Mismatches 3; Indels
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COMPUTER RADABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb
COMPUTER: IBM compatible PC
COMPUTER: IBM compatible PC
COMPUTER: Word 97 (DOS text format)
CURRENT APPLICATION NUMBER: US/09/784,423
FILING DATE: 15-Feb-2001
CLASSIFICATION: «Unknown»
                                                                                                                                                                                                                                                                                                                                                                                                                                       Bacher, Jeffery W.
TITLE OF INVENTION: MATERIALS AND METHODS FOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.8%; Score 18.2; DB 1;
Best Local Similarity 87.0%; Pred. No. 4e+02;
Matches 20; Conservative 0; Mismatches 3;
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APPLICATION NUMBER: 09/018,584
FILING DATE: 04-Feb-1998
ATTORNEY/AGENT INFORMATION:
NAME: Grady J. Frenchick
REGISTRATION NUMBER: 29,018
REFERENCE/DOCKET NUMBER: 16026.9180
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REPEAT DNA MARKERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Promega Corporation
STREET: 2800 Woods Hollow Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: Linear
SEQUENCE DESCRIPTION: SEQ ID NO: 96
                                                                                                                                                                                 2117 TGTTACCCAGGCTGGAGTGCAGT 2139
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                                                                                                                                                                                                                            23 TATCACCCAGGCTGGAGTGCAAT 1
                                                                                                                                                                                                                                                                                                                                          Sequence 96, Application US/09784423
Patent No. 6767703
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 96
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: Schumm, James W.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CITY: Madison
STRANDEDNESS: Sir
TOPOLOGY: Linear
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                     ; TOPOLOGY:
US-09-018-584A-96
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USA
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APPLICANT: Alnemri, Emad S.
APPLICANT: Fernandez-Alnemri, Teresa
APPLICANT: Fernandez-Alnemri, Teresa
APPLICANT: Fernandez-Alnemri, Teresa
TITLE OF INVENTION: USING THE SAME, AND COMPOSITIONS FOR AND METHODS
TITLE OF INVENTION: OF MAKING THE SAME
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcocck, Washburn, Kurtz, Mackiewicz & No. 6037461ris
GITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
OTHER INFORMATION: Description of Artificial Sequence:Synthetic US-09-156-253-30
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             APPLICANT: Montal Brett P.
APPLICANT: Montal Brett P.
APPLICANT: Baker, Brenda F.
APPLICANT: Baker, Brenda F.
APPLICANT: Cowsert, Lex M.
TITLE OF INVENTION: Antisense Modulation of CREL Expression FILE REFERENCE: RTS-0010
CURRENT PELLING DATE: 1998-09-18
NUMBER OF SEQ ID NOS: 48
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 30
LENGTH: 18
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100.0%; Pred. No. 4.4e+02;
ive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                         0.8%; Score 18; DB 1; Length 18;
100.0%; Pred. No. 4.4e+02;
tive 0; Mismatches 0; Indels
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ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: WINDOWS
SOFTWARE: WordPerfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/859,167
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FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: DeLuca, Mark
REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: TUU-
TELECOMMUNICATION INFORMATION:
TELEPAX: (215) 568-3100
TELEPAX: (215) 568-3100
TELEPAX: (215) 568-3100
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
LENGTH: 18 base pairs
LENGTH: 18 base pairs
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Patent No. 6037461
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Best Local Similarity 100.
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity 100.
Matches 18; Conservative
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: DNA
US-08-859-167-9
GENERAL INFORMATION:
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STATE:
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9y 2334 GCCTCCCAAAGTGCTGG 18

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Us-19-13-273-9

Sequence 9, Application Us/0910973

Faterin 80. 6663760

GREBAL INCOMPATION:
APPLICANT: Allenti, Enad 8.
APPLICANT: Allenti, Enad 8.
TITLE OF INVENTION: GC MALLE FORE
TITLE OF INVENTION: GC MALLE FORE
APPLICANT: BLIASE, IT COMPATION: GC MALLE FORE
APPLICANT MALLE FORE
APPLICANT AND APPLICATION AND APPLICATION AND APPLICATION APPLICATION AND APPLICANT APPLICANT AND APPLICANT APPLICANT AND APPLICANT APPLICANT APPLICANT AND APPLICANT A
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GENERAL INFORMATION:
APPLICANT: DePinho, Robert A.
TITLE OF INVENTION: A METHOD OF INHIBITING CELL PROLIFERATION USING AN ANTI-ONCOGENE
FILE REPERENCE: 96700/469
CURRENT APPLICATION NUMBER: US/09/044,602
CURRENT PILING DATE: 1998-03-19
NUMBER OF SEQ ID NOS: 2
SOFTWARE: Patentin version 3.1
SOFTWARE: Patentin version 3.1
LENGTH: 18
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APPLICANT: Baird, Stephen
APPLICANT: Baird, Stephen
APPLICANT: Holoik, Martin
APPLICANT: Young, Sean
TITLE OF INVENTION: Antisense IAP Nucleic Acids and Uses
TITLE OF INVENTION: Thereof
FILE REFERENCE: 07891/02501
CURRENT FILING DATE: 2000-09-28
NUMBER OF SEQ ID NOS: 231
SOFTWARE: FastSEQ for Windows Version 4.0
SEQ ID NO 98
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.8%; Score 18; DB 1; Length 18; Best Local Similarity 100.0%; Pred. No. 4.4e+02; Matches 18; Conservative 0; Mismatches 0; Indels
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0.8%; Score 18; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 4.4e+02;
Matches 18; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: primer for MDM2 mutant
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:

OTHER INFORMATION: based on Homo sapiens US-09-672-717-98
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-780-173A-18/c
; Sequence 18, Application US/09780173A
Patent No. 6455307
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 98, Application US/09672717; Patent No. 6673917
GENERAL INFORMATION:
APPLICANT: Korneluk, Robert G.
                                                                                                 2334 GGCCTCCCAAAGTGCTGG 2351
                                                                                                                                                                                                                                                       ; Sequence 2, Application US/09044602
; Patent No. 6613750
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1271 CAGATGTTGGGCCCTTCG 1288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2189 TCTCCTGCCTCAGCCTCC 2206
                                                                                                                           1 GGCCTCCCAAAGTGCTGG 18
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-672-717-98/c
                                                                                                                                                                                                               RESULT 516
US-09-044-602-2/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-044-602-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:
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Sequence 9, Application US/09723450
Patent No. 6556751
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL INFORMATION:
GENERAL TAILE OF INVENTION: Compositions For And Methods Of Making The Same, ATITIES OF INVENTION: Compositions For And Methods Of Making The Same
FILE REFERENCE: TJU12445
CURRENT APPLICATION NUMBER: US/09/723,450
CURRENT APPLICATION NUMBER: 09/276,993
PRIOR FILING DATE: 1999-03-26
PRIOR FILING DATE: 1999-03-26
PRIOR FILING DATE: 1999-05-20
NUMBER OF SEQ ID NOS: 17
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STREET: Woodcock, Washburn, Kurtz, Mackiewicz & No. 6207801ris STREET: One Liberty Place, 46th floor CITY: Philadelphia STATE: Parts: USA COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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; OTHER INFORMATION: No. 6576751el Sequence
US-09-723-450-9
                                                                                                                                                                                                                                                  MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: WINDOWS
SOFTWARE: WORDFORTEL
CURRENT APPLICATION NUMBER: US/09/276,993
                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/859,167
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: DeLuca, Mark
REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: TJU-
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2334 GGCCTCCCAAAGTGCTGG 2351
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 9: SEQUENCE CHARACTERISTICS: LENGTH: 18 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
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US-09-723-450-9
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LENGTH: 18
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                                                                                                                          COMPUTER: FELORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Flam PC compatible
COMPUTER: PatentIn Felease #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/097,199
FILING DATE:
CLASSIFICATION:
PRICH APPLICATION:
APPLICATION NUMBER: US 08/692,787
FILING DATE:
APPLICATION NUMBER: US 08/692,787
FILING DATE:
APPLICATION NUMBER: P-42,023
REFERENCE/DOCKET NUMBER: URCC:018
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEFRAK: (512) 414-3000
TELEFRAK: (512) 414-7577
INFORMATION FOR SEQ ID NO: 87:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COFTWARE PARENTIN STATEM: C-DOS/MS-DOS SOFTWARE PARENTIN Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
APPLICATION UNDRER: US/08/133,629
FILING DATE: 07-0CT-1993
CLASSIFICATION: 435
ATTONENY/AGENT INFORMATION:
NAME: Greer, Helen
REGISTRATION NUMBER: 36,816
REGISTRATION NUMBER: 36,816
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
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Patent No. 5597694
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT:
MUNICO, David E.
TITLE OF INVENTION: AMPLIFICATION OF NUCLEIC ACIDS NUMBER OF SEQUENCES: 8
CORRESPONDENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSE: Wolf, Greenfield & Sacks, P.C.
STREET: 600 Atlantic Avenue
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Boston
STATE: Massachusetts
COUNTX: United States of America
ZIP: 02210
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2336 CCTCCCAAAGTGCTGGGA 2353
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            22 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; TOPOLOGY: linear
US-09-097-199-87
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 521
US-08-133-629-2/c
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...culCANT: Susan M Freier;
APPLICANT: Susan M Freier;
TITLE OF INVENTION: ANTISENSE MODULATION OF CASEIN KINASE 2-ALPHA PRIME EXPRESSION;
FILE REFERENCE: RTS-0165
CURRENT APPLICATION NUMBER: US/09/780,173A
CURRENT FILING DATE: 2001-02-08
NUMBER OF SEQ ID NOS: 95
LENGTH: 20
TYPE.
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Patent No. 6566135

GENERAL INFORMATION:

APPLICANT: Vickie L. Brown-Driver

APPLICANT: Andrew T. Watt

TITLE OF INVENTION: ANTISENSE MODULATION OF CASPASE 6 EXPRESSION

TITLE OF INVENTION NUMBER: US/09/679,299A

CURRENT FILING DATE: 2000-10-04

NUMBER OF SEQ ID NOS: 164

LENGTH: 20
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Requence 87, Application US/09097199

Retent No. 6218529

GENERAL INFORMATION:

APPLICANT: 0'Hara, S. Mark

APPLICANT: Ralph, David

APPLICANT: Neltri, Robert

TITLE OF INVENTION: BIOMARKERS AND TARGETS FOR DIAGNOSIS,

TITLE OF INVENTION: PROGNOSIS AND MANAGEMENT OF PROSTATE DISEASE

NUMBER OF SEQUENCES: 87

CORRESPONDENCE ADDRESS:

ADDRESSEE: Arnold, White & Durkee
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .
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ilarity 100.0%; Pred. No. 4.3e+02;
Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: Antisense Oligonucleotide US-09-780-173A-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2124 CAGGCTGGAGTGCAGTGG 2141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2187 ATTCTCTGCCTCAGCCT 2204
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                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ) OTHER INFORMATION: PCR Primer US-09-679-299A-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STREET: P.O. Box 4433
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Best Local Similarity
Matches 18; Conserva
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US-09-679-299A-4/c
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Fatent No. 6043088
GENERAL INFORMATION:
APPLICANT: Bookstein, Robert
APPLICANT: Isaacs, William B.
TITLE OF INVENTION: Gene Located on Human Chromosome 8
ITILE OF INVENTION: Gene Located on Human Chromosome 8
INTRESSEE: Campbell and Flores
CORRESPONDENCE 5.59
CORRESPONDENCE 5.69
CORRESPONDENCE ADDRESS:
ADDRESSEE: Campbell and Flores
STREET: 4370 La Jolla Village Drive, Suite 700
CITY: San Diego
STATE: California
COUNTRY: USA
ZIP: 92122
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION NUMBER: US/08/445,515
                                                                                                                                                                           Gaps
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                                                                                                                                                                                                                                                                                                                                                                                Sequence 15, Application US/08933358
Patent No. 6013444
GENERAL INFORMATION:
APPLICANT: Dau, Peter C.
APPLICANT: Liu, Debang
TITLE OF INVENTION: DIA BRACKETING LOCUS COMPATIBLE STANDARDS FOR
FILE REFERENCE: 4340018
FILE REFERENCE: 4340018
CURRENT APPLICATION NUMBER: US/08/933,358
CURRENT FILING DATE: 1997-09-18
NUMBER OF SEQ. ID NOS: 2.0
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                                                                                                                    Query Match 0.8%; Score 17.8; DB 1; Length 21; Best Local Similarity 90.5%; Pred. No. 4.3e+02; Matches 19; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
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CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn
                           single
TYPE: nucleic acid
STRANDEDNESS: sing
                                               ; TOPOLOGY: linear
US-08-133-629-2
                                                                                                                                                                                                                                                                                                                                                                      US-08-933-358-15/c
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LENGTH: 21
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                                                                                                                                                                                                                                                          Length 21;
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                                                                                                                                                                                                                                                                                                    2; Indels
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Identification and Banding of
Specific Human Chromosomes and
Regions
                                                                                                                                                                                                                                                          0.8%; Score 17.8; DB 1;
90.5%; Pred. No. 4.3e+02;
tive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.8%; Score 17.8; DB 1; Best Local Similarity 90.5%; Pred. No. 4.3e+02; Matches 19; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 87, Application US/09918686
Fatent No. 6475739
GENERAL INFORMATION:
APPLICANT: Brunkow, Mary
APPLICANT: Breaching-Hampton, Karen
APPLICANT: Seaching-Hampton, Karen
TITLE OF INVENTION: GENOMIC DELETIONS
TITLE OF INVENTION: GENOMIC DELETIONS
FILE REFERENCE: 240083.515
CURRENT APPLICATION NUMBER: US/09/918,686
CURRENT FILING DATE: 2001-07-30
NUMBER OF SEQ ID NOS: 105
SEQ ID NOS: 105
LENGTH: 21
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US-08-070-517-2/C
US-08-070-517-2/C
Sequence 2, Application US/08070517
Patent No. 5538869
GENERAL INFORMATION:
APPLICANT: Michael J. Siciliano
APPLICANT: Pu Liu
TITLE OF INVENTION: In-Situ Hybridizati;
TITLE OF INVENTION: Identification and
TITLE OF INVENTION: Specific Human Chro,
TITLE OF INVENTION: Regions
NUMBER OF SEQUENCES: 5
CORRESPONDENCE ADDRESS:
STREET: P.O. BOX 4433
              REFERENCE/DOCKET NUMBER: P-CJ 1607
TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEFAX: (619) 535-9949
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                         2107 AGTOTTGCTCTGTTACCCAGG 2127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2312 GACCTCGTGATCCGCCCACCT 2332
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31,815
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: PCR primer
                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 90.5
Matches 19; Conservative
                                                                                                                                  LENGTH: 21 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
REGISTRATION NUMBER:
                                                                                                                                                                                              ; TOPOLOGY: linear
US-08-445-515-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Houston
                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-918-686-87/c
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STATE: Te
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-918-686-87
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MOLECULE TYPE: DNA (genomic) HYPOTHETICAL: NO
                                                                                                              Query Match
Best Local Similarity 94.4%;
Matches 17; Conservative
                                                                                                                                                                                                                                           19 CAGGCTGGAGTGCARTGG
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CITY: Lexington
STATE: Massachusetts
linear
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TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
                                                                   US-08-118-441-2
                                                                                                                                                                                                                                                                                                                                       RESULT 527
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| Sequence 2, Application US/08118441 |
| Patent No. 5578493 |
| Patent No. 5578493 |
| GENERAL INFORMATION: |
| APPLICANT: G11liam, T. Conrad |
| APPLICANT: G11liam, T. Conrad |
| TITLE OF INVENTION: ISOLATION AND USES OF A WILSON'S DISEASE |
| TITLE OF INVENTION: GENE |
| NUMBER OF SEQUENCES: 29 |
| CORRESPONDENCE ADDRESS: |
| ADDRESSEE: Cooper & Dunham |
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.7%; Score 17.6; DB 1; Length 19; Best Local Similarity 94.4%; Pred. No. 4.6e+02; Matches 17; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESTONDENESS:
ADDRESSER: Cooper & Lunham
STREET: 30 Rockefeller Plaza
CITY: New York
STATE: New York
COUNTRY: United States of America
ZIP: 10112
COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/NS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/118,441
FILING DATE:
CLASSIFICATION:
NAME: White, John P.
REGISTERATION NUMBER: 28,678
REGISTERATION NUMBER: 28,678
REGISTERATION NUMBER: 0575/44011
TELECOMMUNICATION NUMBER: 0575/44011
CALP: //TALP: //TALP: CAMPUTER READABLE FORM:
MEDIUM TYPE: Floppy Disk
COMPUTER: TBM FC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII-DOS
CURRENT APPLICATION DATH:
APPLICATION NUMBER: US/08/070,517
FILING DATE: 19930601
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: BATBATE S. Kitchell
REGISTRATION NUMBER: 33,928
REFERENCE/DOCKET NUMBER: UTSC:290/KIT
TELECOMMUNICATION INFORMATION:
TELEPAN: (512) 474-7577
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE GIARACTERISTICS:
LENGTH: 19 base pairs
TYPE: NUCLEIC acids
TYPE: NUCLEIC ACIDS
MONOROWENESS: SINGLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2124 CAGGCTGGAGTGCAGTGG 2141
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TELEFAX: (212) 664-0525
TELEX: 42253 COOP UI
INFORMATION FOR SEG ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 19 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-070-517-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ઠે
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WORDSTORY LINE
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DNA (genomic)
           MOLECULE TYPE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COUNTRY: UN
                       ;
US-08-422-706B-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-338-579A-2
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                                                                                                                                            Query Match 0.7%; Score 17.6; DB 1; Length 19; Best Local Similarity 94.4%; Pred. No. 4.6e+02; Matches 17; Conservative 1; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Brook, J. David
APPLICANT: Brook, J. David
APPLICANT: Braw, Duncan J.
APPLICANT: Shaw, Duncan J.
APPLICANT: Ariely, Helen G.
APPLICANT: Johnson, Keith J.
TITLE OF INVENTION: DNA SEQUENCE ENCODING THE MYOTONIC
TITLE OF INVENTION: DYSTROFHY GENE AND USES THEREOF
NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
ADDRESSEE: AMBLICON, Brook, Smith & Reynolds, P.C.
STREET: Two Militia Drive
CITY: Leatington
STATE: Massachusetts
COUNTRY: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
FLING DATE:
APPLICATION NUMBER: US/08/422,706B
FLING APPLICATION NUMBER: US/08/4543
FLING DATE: 08-AUG-1994
FRIOR APPLICATION NUMBER: US/08/23,612
FRICK APPLICATION NUMBER: US/08/23,612
FRICK APPLICATION NUMBER: US/08/23,612
FRICK APPLICATION NUMBER: US/07/839,255
FRICK APPLICATION NUMBER: US/07/839,255
FRICK APPLICATION NUMBER: US/07/839,01545
FILING DATE: 20-FEB-1993
FRICK APPLICATION NUMBER: US/08/31/01545
FILING DATE: US-FEB-1993
FRICK APPLICATION NUMBER: US/08/31/0253
FRICK APPLICATION NUMBER: US/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/31/08/
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REGISTRATION NUMBER: 32,227
REFERENCE/DOCKET NUMBER: MIT-5830A2
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-861-6240
TELEFAX: 617-861-9540
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-08-422-706B-14/c; Sequence 14, Application US/08422706B Patent No. 5977333; GENERAL INPORMATION:
                                                                                                                                                                                                                                                                                                    2124 CAGGCTGGAGTGCAGTGG 2141
                               MOLECULE TYPE: DNA (genomic) US-08-422-699A-14
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: 617-861-9540
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                           19 CAGGCTGCAGTGCARTGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 19 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   02713
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    Length 19;
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                                                                                                                                                                                            RESULT 529
US-08-130-579A-2/C
| Sequence 2, Application US/08338579A
| Patent No. 6068975
| GENERAL INFORMATION:
| APPLICANT: Tania, Rudolph E. TITLE OF INVENTION: ISOLATION AND USES OF A WILSON'S TITLE OF INVENTION: DISEASE GENE NUMBER OF SEQUENCES: 107
| CORRESPONDENCE ADDRESS: ADDRESSE: Cooper & Dunham STREET: 1185 Avenue of the Americas
                                              0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 2, Application PC/TUS9409851
Sequence 2, Application PC/TUS9409851
GENERAL INFORMATION:
APPLICANT: Gilliam, T. Conrad
APPLICANT: Tanzai, Rudolph E.
TITLE OF INVENTION: ISOLATION AND USES OF A WILSON'S
TITLE OF INVENTION: DISEASE GENE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER FRADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/38,579A
FILING DATE: June 17, 1996
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: White, John P.
REGISTRATION NUMBER: 28,678
REFERENCE/DOCKET NUMBER: 0575/44011-A-PCT-US
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match

O.7%; Score 17.6; DB 1;
Best Local Similarity 94.4%; Pred. No. 4.6e+02;
Matches 17; Conservative 1; Mismatches 0;
Score 17.6; DB 1;
Pred. No. 4.6e+02;
1; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   New York
: United States of America
                                                                                     2124 CAGGCTGGAGTGCAGTGG 2141
                                                                                                                 CAGGCTGGAGTGCARTGG 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEPHONE: (212) 278-0400
TELEFAX: (212) 391-0525
Query Match 0.7%;
Best Local Similarity 94.4%;
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2124 CAGGCTGGAGTGCAGTGG
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  linear
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NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REPERENCE/DOCKET NUMBER: 09865.601
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 831-2106
TELEFAX: (608) 831-2106
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-09-564-805-100/c
; Sequence 100, Application US/09564805
; Patent No. 6333403
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                             2331 CTCGGCCTCCCAAAGTGCT 2349
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2196 CCTCAGCCTCCCAATTAGC 2214
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                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
INMEDIATE SOURCE:
CLONE: mfd103p2
US-08-222-177A-330
                                                                                                             TELEX:
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Homo sapiens
US-09-564-805-100
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US-08-222-177A-330
is Sequence 330. Application US/08222177A
is Patent No. 5582979
is GENERAL INFORMATION:
INTER OF INVENTION: LENGTH POLYMORPHISMS IN
TITLE OF INVENTION: (dC-dA)n.(dG-dT)n SEQUENCES AND METHODS OF USING SAME
INVERSE OF SEQUENCES: 460
CORRESPONDENCE ADDRESS:
ADDRESSE: Dewitt Ross & Stevens, S.C.
STREET: 8000 Excleior Drive, Suite 401
CORRESPONDENCE ADDRESS:
STREET: Madison
COUNTRY: USA
STATE: Wisconsin
COUNTRY: READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: Depthy disk
COMPUTER: Patentin Release #1.0, Version #1.25
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Pred. No. 4.6e+02;
1; Mismatches 0; Indels
                                                                 CITY: New York
STATE: New York
COUNTRY: United States of America
ZIP: 10112
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: ISM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US94/09851
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                                      CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2124 CAGGCTGGAGTGCAGTGG 2141
                 CORRESPONDENCE ADDRESS:
ADDRESSEE: Cooper & Dunham
STREET: 30 Rockefeller Plaza
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    19 CAGGCTGGAGTGCARTGG 2
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Best Local Similarity 94.44
Matches 17; Conservative
NUMBER OF SEQUENCES: 92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: DN HYPOTHETICAL: NO
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APPLICANT: Tavidjan, Sean V.
APPLICANT: Tavidjan, Sean V.
APPLICANT: Tavidjan, Sean V.
APPLICANT: Teng, David H.F.
APPLICANT: Simarci, Jacques
APPLICANT: Rommens, Johanna M.
APPLICANT: Myriad Genetics, Inc.
TITLE OF INVENTION: Chromosome 17p-Linked Prostate Cancer Susceptibility
TITLE OF INVENTION: Gene and a Paralog and Orthologous Genes
TITLE OF INVENTION: Gene and a Paralog and Orthologous Genes
FILE REFERENCE: 2318-258
CURRENT APPLICATION NUMBER: US/09/564,805
FRIOR PAPLICATION NUMBER: US 60/107,468
PRIOR APPLICATION NUMBER: 09/434,382
PRIOR FILING DATE: 1999-11-05
NUMBER OF SEQ ID NOS: 240
SEQ ID NO 100
SEQ ID NO 100
SEQ ID NO 100
SEQ ID NO 100
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Query Match

0.7%; Score 17.4; DB 1; Length 19;
Best Local Similarity 94.7%; Pred. No. 4.7e+02;
Matches 18; Conservative 0; Mismatches 1; Indels
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US-09-091-952A-86/c
; Sequence 86, Application US/09091952A
; Patent No. 645852;
; GENERAL INFORMATION:
; APPLICANT: Detera-Wadleigh, Sevilla D. Gershon, Elliot S. Badner, Judith A. Goldin, Lynn R. Berrettini, Wade H.
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PRIOR FILING DATE: 1998-09-25
PRIOR APPLICATION NUMBER: PCT/US99/22283
PRIOR FILING DATE: 1999-09-24
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: double
                                                                                                                                                                   , ORGANISM: Homo Sapiens
US-09-404-912-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CITY: Madison
STATE: Wisconsin
COUNTRY: USA
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US-08-222-177A-351
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MOLECULE TYPE: DI
IMMEDIATE SOURCE:
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                                                                                                                                                TYPE: DNA
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                 Sanders, Alan R.
Esterling, Lisa E.
TITLE OF INVENTION: Chromosomal Markers and Diagnostic Tests for Manic-Depressive Illness
                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
                                                                                                                                                                                                                        COMPUTER READABLE PORM:

COMPUTER READABLE PORM:

MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
COERATING SYSTEM: DOS
SOFTWARE: FASEREG for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/091,952A
FILING DATE: 19-Apr-1999
CLASSIFICATION NUMBER: US 60/029,278
FILING DATE: 28-OCT-1996
APPLICATION NUMBER: PCT/US97/19381
FILING DATE: 28-OCT-1996
APPLICATION NUMBER: PCT/US97/19381
FILING DATE: 28-OCT-1997
ATTORNEY/AGENT INFORMATION:
RAFERENCE/DOCKET NUMBER: 35,367
REFERRATION NUMBER: 35,367
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 3, Application US/09404912
Fatent No. 6703228
GENERAL INFORMATION:
APPLICANT: David Houseman
APPLICANT: Barbara Jordan
APPLICANT: Alain Charest
TITLE OF INVENTION: Genotyping and DNA Analysis
FILE REFERENCE: No656/7045 (HCL/MAT)
CURRENT APPLICATION WUMBER: US/09/404,912
CURRENT FILING DATE: 1999-09-24
PRIOR APPLICATION NUMBER: US 60/101,757
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LOCATION: 1...19
OTHER INFORMATION: D18S378 forward primer SEQUENCE DESCRIPTION: SEQ ID NO: 86:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (415) 576-0200
TELEFAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2111 TTGCTCTGTTACCCAGGCT 2129
Yoshikawa, Takec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  INFORMATION FOR SEQ ID NO: 86:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    rrecrerereacecadeer 1
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STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity 94.7
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME/KEY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 534
US-09-404-912-3/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-09-091-952A-86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE
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US-08-222-1774-351/c

J Sequence 351, Application US/08222177A

Sequence 351, Application US/08222177A

Sequence 351, Application US/08222177A

Sequence 351, Application US/08222177A

J THORNATION: GENERAL INFORMATION: Langth POLYMORPHISMS IN

TITLE OF INVENTION: (dG-dA)n.(dG-dT)n.SEQUENCES AND METHODS OF USING SAME

NUMBER OF SEQUENCES: 460

CORRESPONDENCE ADDRESSE: Dewlitt Ross & Stevens, S.C.

STREET: 8000 Excelsior Drive, Suite 401
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                                                                                                                                                                                                                                               Length 19;
                                                                                                                                                                                                                                                                                                       1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/222,177A
                                                                                                                                                                                                                                            Query Match

0.7%; Score 17.4; DB 1;
Best Local Similarity 94.7%; Pred. No. 4.7e+02;
Matches 18; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/341,562
FILING DATE: 21-APR-1989
ATTORNEY/AGENT INFORMATION:
NAME: SAZA, CHAILGS S.
REFERENCE/DOCKET NUMBER: 09865.601
TELECHMUNICATION INFORMATION:
NUMBER OF SEQ ID NOS: 691
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 3
LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: USA
ZIP: 53717-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                  2144 GATCTTGGCTCACTGCAAG 2162
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Gaps
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STREET: 1100 No. 6555654th Glebe Road, Eighth Floor
                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPATION SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/060,299
FILING DATE: 15-APR-1998
CLASSIFICATION NUMBER: US 60/043,553
FILING DATE: 15-APR-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 15-APR-1997
ATTONEY/AGENT INFORMATION:
NAME: B.J.Sadoff
REGISTRATION NUMBER: 36,663
REGISTRATION NUMBER: 36,663
FILING DATE: (703)816-4091
TELEPHONE: (703)816-4091
TELEPHONE: (703)816-4091
TELEPHONE: (703)816-4091
TELEPHONE: AD DASE PAIRE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 17.4; DB 1; Length 20;
Pred. No. 4.6e+02;
0; Mismatches 1; Indels
                                                               APPLICANT: May, Fall Lines APPLICANT: Marinan, Tony R APPLICANT: Merrinan, Tony R APPLICANT: Merrinan, Tony R APPLICANT: Metzker, Michael L TITLE OF INVENTION: No. 6545137el Receptor NUMBER OF SEQUENCES: 455 CORRESPONDENCE ADDRESSE: ADDRESSE: ADDRESSE: ADDRESSE: ADDRESSE: ADDRESSE: ADDRESSE: Uson and Vanderhye STREET: 1100 No. 6545137th Glebe Road, Eighth Floor CITY: Arlington STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hey, Patricia
Kawaguchi, Yoshihiko
Merziman, Tony R
Metzker, Michael L
TITLE OF INVENTION: NO. 6555654el LDL-Receptor
NUMBER OF SEQUENCES: 455
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 286, Application US/09402923A
Patent No. 655654
GENERAL INFORMATION:
APPLICANT: Todd, John A
Hess, John W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Caskey, Charles T
Cox, Roger D
Gerhold, David
Hammond, Holly
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  Gerhold, David
Hammond, Holly
Hey, Patricia
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Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-402-923A-286
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US-09-060-299-286
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                                                                                   Sequence 18, Application US/09851896

Fatent No. 6410325

GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: State Matt
TITLE OF INVENTION: ANTISENSE MODULATION OF PHOSPHOLIPASE A2, GROUP VI (CA2+-INDEPEND FILE REFERENCE: RTS-0220
CURRENT APPLICATION NUMBER: US/09/851,896
CURRENT FILING DATE: 2001-05-08
NUMBER OF SEQ ID NOS: 89
SEQ ID NO 18
LENGTH: 20
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Sequence 95, Application US/09898361

Parent No. 6603152

GENERAL INFORMATION:
APPLICANT: Susan Murray
APPLICANT: Jacquelline Wyatt
TITLE OF INVENTION: EXPRESSION
TITLE OF INVENTION: EXPRESSION
TITLE OF INVENTION NUMBER: US/09/898,361

CURRENT APPLICATION NUMBER: US/09/898,361

CURRENT FILING DATE: 2001-06-21

NUMBER OF SEQ ID NOS: 163

SEQ ID NO 95

LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.7%; Score 17.4; DB 1; Length 20; Best Local Similarity 94.7%; Pred. No. 4.6e+02; Matches 18; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.7%; Score 17.4; DB 1; Length 20; Best Local Similarity 94.7%; Pred. No. 4.6e+02; Matches 18; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ) OTHER INFORMATION: Antisense Oligonucleotide US-09-898-361-95
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Antisense Oligonucleotide US-09-851-896-18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 286, Application US/09060299
Patent No. 6545137
GENERAL INFORMATION:
APPLICANT: Todd, John A
APPLICANT: Hess, John W
APPLICANT: Caskey, Charles T
APPLICANT: Coskey, Charles T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2334 GGCCTCCCAAAGTGCTGGG 2352
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19 CCACAACACCTGGCTAATT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-060-299-286
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                                                                                        US-09-851-896-18
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0.7%; Score 17.4; DB 1; Length 20; llarity 94.7%; Pred. No. 4.6e+02; Conservative 0; Mismatches 1; Indels
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Sequence 28, Application US/08781891
Sequence 28, Application US/08781891
Sequence 28, Application US/08781891
Sequence 28, Application US/08781891
September 10, Ving-Hui
APPLICANT: Yu, Chang-En
APPLICANT: Wulligan, John T.
APPLICANT: Schellenberg, Gerald D.
TITLE OF INVENTION: GENE AND GENE PRODUCTS RELATED TO
TITLE OF INVENTION: WERNER'S SYNDROME
NUMBER OF SEQUENCES: 209
CORRESPONDENCE ADDRESS:
ADDRESSEE: SEED and BERRY LLP
STREET: 6300 Columbia Center, 701 Fifth Avenue
                                                                                                                       US-09-544-39BB-582/C
; Sequence 582, Application US/09544398B
; Patent No. 6770461
; GENERAL INFORMATION;
; APPLICANT: Carulli, John P.
; APPLICANT: Little, Randall D.
; APPLICANT: Little, Randall D.
; APPLICANT: Acker, Robert R.
; APPLICANT: Recker, Robert R.
; TITLE OF INVENTION: High bone mass gene of 11q13.3
; TITLE OF INVENTION: High bone mass gene of 11q13.3
; TITLE OF INVENTION WIMBER: US/09/544,398B
; CURRENT APPLICATION WIMBER: US 09/229,319
; PRIOR APPLICATION WIMBER: US 09/229,319
; PRIOR FILING DATE: 1999-01-13
; PRIOR FILING DATE: 1999-10-13
; PRIOR FILING DATE: 1999-10-23
; NUMBER OF SEQ ID NOS: 641
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO S: LEMMIN DATE: DAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Floppy disk
COMPUTER: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/781,891
CLASSIFICATION: 800
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NAME: No. 6090620tenburg Ph.D., Carol
REGISTRATION NUMBER: 39,317
REFERENCE/DOCKET NUMBER: 240052.419
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GTTACCC 2124
          1 GCTGGGATTACAGGCGTGA 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-544-398B-582
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2106 GAGICTIGCICI
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STATE: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
nes 18; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Loca
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 542
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         셤
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Sequence 30, Application US/09574779B

Batent No. 6/67720

GENERAL INFORMATION

TITLE OF INVENTION: No. 6/67720el cDNAs encoding catenin-binding proteins with TITLE OF INVENTION: Encoding and/or gene regulation FILE REFERENCE: 26/6-4415US

CURRENT APPLICATION NUMBER: US/09/574,779B

CURRENT PILING DATE: 2000-05-19

PRIOR APPLICATION NUMBER: 99201543.8

PRIOR FILING DATE: 1999-05-17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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US-09-574-779B-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
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                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
0.7%; Score 17.4; DB 1; Length 20;
Best Local Similarity 94.7%; Pred. No. 4.6e+02;
Matches 18; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.7%; Score 17.4; DB 1; Length 20; Best Local Similarity 94.7%; Pred. No. 4.6e+02; Matches 18; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: PCT/GB98/01102
FILING DATE: 15-APR-1998
APPLICATION NUMBER: US 60/043,553
FILING DATE: 15-APR-1997
APPLICATION NUMBER: US 60/048,740
FILING DATE: 05-JUN-1997
ATTORNEY AGENT INFORMATION:
NAME: B.J.Sadoff
                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: US/09/402,923A FILING DATE: 14-Feb-2001 PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 286:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REGISTRATION NUMBER: 36,663
REFERENCE/DOCKET NUMBER: 620-81
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2146 TCTTGGCTCACTGCAAGCT 2164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (703)816-4091
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (703)816-4100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 286: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
CITY: Arlington
STATE: Virginia
COUNTRY: US
ZIP: VA 22201-4714
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 158
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 30
LENGTH: 20
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2347 GCTGGGATTACAGGCATGA 2365

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TELEFAX:
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                                                                                                                                                                                                      0.7%; Score 17.4; DB 1; Length 21; 94.7%; Pred. No. 4.6e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.7%; Score 17.4; DB 1; Length 21; 94.7%; Pred. No. 4.6e+02; ive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTEY USA

ZIP: 98104-7092

ZIP: 98104-7092

COMPUTER READABLE FORD:

MEDIUM TYPE: Floppy disk

COMPUTER: IBM PC compatible

OPERATING SYTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/618,166

FILING DATE: 17-Jul-2000

CLASSIFICATION: <UNknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mulligan, John T.
Schellenberg, Gerald D.
TITLE OF INVENTION: GENE AND GENE PRODUCTS RELATED TO
WERNER'S SYNDROME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CORRESPONDENCE ADDRESS:
ADDRESSEE: Seed Intellectual Property Law Group
STREET: 701 Fifth Avenue, Suite 6300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: MCMABLETS, DAVID D.
REGISTRATION NUMBER: 33,963
REFERENCE/DOCKET NUMBER: 240052.419C1
TELEPHONE: (206) 622-4900
TELEPRAK: (206) 622-4900
TELEPRAK: (206) 622-4900
TELEPRAK: (206) 622-6031
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE DESCRIPTION: SEQ ID NO: 28:
US-09-618-166-28
                                                                                                                                                                                                                                                                                                                                                                                        RESULT 343
US-09-618-166-28/C
; Sequence 28, Application US/09618166
; Patent No. 6583112
; GENERAL INFORMATION:
; APPLICANT: Fu, Ying-Hui
; APPLICANT: Fu, Ying-Hui
; Junko
                                                                                                                                                                                                                                                                                       2294 GGATGGTCTCGATCTCCTG 2312
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                TELEFAX: (206) 682-6031
INFORMATION FOR SEQ ID NO: 28:
SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
TYPE: nucleic acid
STRANBEDNESS: single
                                                                                                                                                                                                                                                                                                                               21 GGATGGTCTCGAACTCCTG 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE CHARACTERISTICS:
LENGTH: 21 base pairs
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STRANDEDNESS: single
  (206) 622-4900
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CITY: Seattle
STATE: Washington
                                                                                                                                                                                                    Query Match
Best Local Similarity 94.7
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear
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Best Local Similarity 94.7°
Matches 18; Conservative
                                                                                                                                           ; TOPOLOGY: linear
US-08-781-891-28
TELEPHONE:
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| Sequence 1, Application US/08837302
| Patent No. 568741
| GENERAL INFORMATION: Perv, Scott E. APPLICANT: Targan, Stephan R. TITLE OF INVENTION: Methods of Diagnosing a Medically TITLE OF INVENTION: Resistant Clinical Subtype of Ulcerative Colitis NUMBER OF SEQUENCES: 10
| CORRESPONDENCE ADDRESS: Campbell & Flores Lip STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 .
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRAILING SISIEM: E-LOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRANT APPLICATION DATA:
APPLICATION NUMBER: US/08/133,629
FILING DATE: 07-0CT-1993
CLASSIFICATION: 435
ATTONREY/AGENT INFORMATION:
NAWE: Greer, Helen
REGISTRATION NUMBER: 36,816
REGISTRATION NUMBER: 36,816
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION:
US-08-133-629-4/c

Sequence 4, Application US/08133629

Patent No. 5597694

GENERAL INFORMATION:

APPLICANT: Munroe, David J.

APPLICANT: Housman, David B.

TITLE OF INVENTION: AMPLIFICATION OF NUCLEIC ACIDS

NUMBER OF SEQUENCES: 8

CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Patentin Release #1.0, Version #1.25
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0.7%; Score 17.2; DB 1;
Best Local Similarity 80.0%; Pred. No. 4.8e+02;
Matches 16; Conservative 3; Mismatches 1;
                                                                                                                                                                                                                        E: Wolf, Greenfield & Sacks,
600 Atlantic Avenue
                                                                                                                                                                                                                                                  STREET: 600 Atlantic Avenue
CITY: Boston
STATE: Massachusetts
COUNTRY: United States of America
                                                                                                                                                                                                                                                                                                                                           ZIP: 02210
COMPUTER REAABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2100 GAGACCGAGTCTTGCTCTGT 2119
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CITY: San Diego
STATE: California
COUNTRY: United States
ZIP: 92122
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEX: 92-1742 EZEKIEL INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               617-720-2441
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TOPOLOGY: linear
US-08-133-629-4
                                                                                                                                                                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-837-302-1/c
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JS-08-798-668-1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 547
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Sequence 1, Application US/08798668

Sequence 1, Application US/08798668

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 90071

COMPUTER READABLE FORM:
MEDIUM TYPER: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION NATR:
APPLICATION NUMBER: US/08/798,668
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION:
APPLICATION UDMBER: US 08/245,297
FILING DATE: 17-MAY-1994
ATTORNEY/AGENT INFORMATION:
NAME: WHITEFORD, WENDY A
REGISTRATION NUMBER: 36,964
REFERENCE/DOCKET NUMBER: P07 32313
TELECOMMUNICATION INFORMATION:
APPLICATION NUMBER: US/08/837,302
FILING DATE: 11-APR-1997
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2115 TCTGTTACCCAGGCTGGAGTGC 2136
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                                     FILING DATE: 11-APR-1997
CLASSIFCATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31,815
REFERENCE/DOCKET NUMBER: 91,815
REFERENCE/DOCKET NUMBER: 9-PM
TELEPHONE: (619) 535-9001
TELEPHONE: (619) 535-9001
TELEPHONE: (619) 535-9001
TELEPHONE: (619) 535-901
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE: (213) 489-4442
TELEFAX: (213) 489-4210
INFORMATION FOR SEQ ID NO: 1.
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 22 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linea
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Gequence 1, Application US/08855825
Fatent No. 6183951
GENERAL INFORMATION:
Targan, Scott E.
Taylor, Kent
Barry, Mary J.
TITLE OF INVENTION: Methods of Diagnosing Clinical Subtypes
of Crohn's Disease with Characteristic Responsiveness to
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  Length 22;
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                                                 Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYDE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ADDRESSEE: Campbell & Flores LLP
STREET: 4370 La Jolla Village Drive, Suite 700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match

0.7%; Score 17.2; DB 1;
Best Local Similarity 86.4%; Pred. No. 4.6e+02;
Matches 19; Conservative 0; Mismatches 3;
Query Match

0.7%; Score 17.2; DB 1;
Best Local Similarity 86.4%; Pred. No. 4.6e+02;
Matches 19; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ATTORNEY/AGENT INFORMATION:
NAME: Campbell, Cathryn A.
REGISTRATION NUMBER: 31, 915
REFERENCE/DOCKET NUMBER: P-PM 2591
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US/08/855,825
FILING DATE: 12-May-1997
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: linear SEQUENCE DESCRIPTION: SEQ ID NO: 1:
                                                                                            2115 TCTGTTACCCAGGCTGGAGTGC 2136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2115 TCTGTTACCCAGGCTGGAGTGC 2136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELECOMMUNICATION INFORMATION:
TELEPHONE: (619) 535-9001
TELEFAX: (619) 535-8949
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
                                                                                                                                         22 réreredéctradecresagnec 1
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US-09-395-345-28/c
Sequence 28 Application US/09395345
Patent No. 6376176
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CITY: San Diego
STATE: California
COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGTH: 22 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Taylor, Kent D. APPLICANT: Rotter, Jerome APPLICANT: Yang, Huiying
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Gaps
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Batent No. 6066459

GENERAL INFORMATION:
APPLICANT: YUVAL GARINI ET AL.
TITLE OF INVENTION: MULTICOLOR CHROMOSOME PAINTING AND TITLE OF INVENTION: MULTICOLOR CHROMOSOME PAINTING AND BANDING NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSEE: Mark M. Friedman c/o Anthony Castorina STREET: 20001 Jefferson Davis Highway, Suite 207
CITY: Alington
STREET: Alington
STREET: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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                                                 COMPUTER READBLE FORM:
MEDIUM TYPE: 1.44 megabyte, 3.5" microdisk
COMPUTER: Twinhead* Slimnote-890TX
OPERATING SYSTEM: MS DOS version 6.2,
OPERATING SYSTEM: Mindows version 3.11
SOFTWARE: Word for Windows version 2.0
SOFTWARE: 22-Apr-1996
FILING DATE: 22-Apr-1996
FILING DATE: 13-Apr-1996
FILING DATE: 18-Aug-93
FILING DATE: 12-Dec-95
APPLICATION NUMBER: 08/571,047
FILING DATE: 21-Feb-95
APPLICATION NUMBER: 08/571,047
FILING DATE: 20-Dec-95
ATTORNEY AGENT INPORMATION:
NAWE: Friedmann, Mark M.
REGISTRATION NUMBER: 33,883
REFREENCE/DOCKET NUMBER: 205/15
TELECOMMUNICATION INPORMATION:
NAWE: FRIEDRATION NUMBER: 205/15
TELECOMMUNICATION INPORMATION:
TELEPHONE: 972-3-562553
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READBLE FORM:
MEDIUM TYPE: 1.44 megabyte, 3.5" microdisk
COMPUTER: Twinhead* Slimnote-890TX
OPERATING SYSTEM: MS DOS version 6.2,
OPERATING SYSTEM: Windows version 3.11
SOFTWARE: Word for Windows version 2.0
SOFTWARE: CONVERTED to ASCI
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/100,104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: Virginia
COUNTRY: United States of America
           United States of America
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2122 CCCAGGCTGGAGTGCAG 2138
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.73
Best Local Similarity 100.0
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 972-3-5625554
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: nucleic acid
STRANDEDNESS: sing]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; TOPOLOGY: linear
US-08-635-820A-2
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US-09-100-104-2/c
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Patent No. 5817462
GENERAL INFORMATION:
APPLICANT: YUVAL GARINI ET AL.
TITLE OF INVENTION: METHOD FOR SIMULTANEOUS DETECTION OF MULTIPLE FLUOROPHORES FOR NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSEE: Mark M. Friedman c/o Robert Sheinbein
STREET: 2940 Birchtree lane
CITY: Silver Spring
STATE: Maryland
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
TITLE OF INVENTION: Methods of Using A Major Histocompatibility Complex TITLE OF INVENTION: Calse III Haplotype To Diagnose Crohn's Disease FILE REFERENCE: P-CE 3639
CURRENT APPLICATION NUMBER: US/09/395,345
CURRENT FILING DATE: 1999-09-13
NUMBER OF SEQ ID NOS: 40
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: BOTTER, JEROME I.
APPLICANT: TOTORA, STOTT E.
APPLICANT: TOTORA, STEPHAN R.
APPLICANT: TOTORA, STEPHAN R.
APPLICANT: TOYORA, HIRROD
APPLICANT: YANG, HUIYING
ITTLE OF INVENTION: WETHOOS OF SCREENING FOR CROHN'S DISEASE
ITTLE OP INVENTION: USING THE MICROSATELLITE ALLELES
ITTLE OP INVENTION: USING THE MICROSATELLITE ALLELES
ITTLE OP INVENTION: 1999-10-15
ITTLE OF INVENTION: UNMBER: 1999-10-15
ITTLE OF INVENTION NUMBER: 08/798,668
RIOR APPLICATION NUMBER: 08/798,668
FRIOR PILING DATE: 1997-02-11
PRIOR APPLICATION NUMBER: 08/245,297
NUMBER OF SEQ ID NOS: 10
SOFTWARE: FRAESEQ for Windows Version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ö
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0.7%; Score 17.2; DB 1; Length 22;
Best Local Similarity 86.4%; Pred. No. 4.6e+02;
Matches 19; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                             3; Indels
                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.7%; Score 17.2; DB 1; Best Local Similarity 86.4%; Pred. No. 4.6e+02; Matches 19; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2115 TCTGTTACCCAGGCTGGAGTGC 2136
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2115 TCTGTTACCCAGGCTGGAGTGC 2136
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 1, Application US/09419406
Patent No. 5334263
GENERAL INFORMATION:
APPLICANT: PLEVY, SCOTT E.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: (genomic) US-09-419-406-1
                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ORGANISM: Homo sapiens
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US-08-635-820A-2/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RESULT 549
US-09-419-406-1/c
                                                                                                                                                                                                                                                                                                            US-09-395-345-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
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US-09-487-445-94/C
; Sequence 94, Application US/09487445
; Patent No. 6258600
; GENERAL INFORMATION:
; APPLICANT: Hong Zhang
; APPLICANT: Lex M. Cowsert
; TITLE OF INVENTION: ANTISENSE MODULATION OF CASPASE 8 EXPRESSION
; FILE REPERBNCE: RTS-0107
; CURRENT FILING DATE: 2000-01-19
; NUMBER OF SEQ ID NOS: 176
; LENGTH: 20
                                                                                                                                                                          Length 20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSE: Anobbe, Martens, Olson & Bear STREET: 620 Newport Center Drive 16th Floor CITY: Newport Beach STATE: 0.8.A. COUNTER: U.S.A. ZIP: 92660 COMPUTER READABLE FORM: MEDIUM TYPE: Diskette COMPUTER: IBM Compatible COMPUTER: IBM Compatible COMPUTER: IBM Compatible COMPUTER: Diskette
                                                                                                                                                                          DB 1; Le
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.7%; Score 17; DB 1; L. Best Local Similarity 100.0%; Pred. No. 4.9e+02; Matches 17; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 12, Application US/08849701
Fatent No. 592254
GENERAL INFORMATION:
APPLICANT: Miyai, Kiyoshi
APPLICANT: Naitoh, Tautomu
APPLICANT: Yonekawa, Toshihiro
ITILE OF INVENTION:
NUMBER OF SEQUENCES: 12
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; FEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-09-487-445-94
                                                                                                                                                                      Query Match 0.7%; Score 17; DB Best Local Similarity 100.0%; Pred. No. 4.5 Matches 17; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SOFTWARE: FastSEQ Version 1.5
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/849,701
FILING DATE:
CLASSIFICATION: 435
                                                                                                                                                                                                                                                      2125 AGGCTGGAGTGCAGTGG 2141
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2125 AGGCTGGAGTGCAGTGG 2141
                  LENGTH: 20 Dage pairs

; TYPE: nucleic acid

; STRANDEDNESS: single

; TOPOLOGY: linear

; MOLECULE TYPE: Genomic DNA

US-09-038-637-155
                                                                                                                                                                                                                                                                                          20 AGCCTGGAGTGCAGTGG 4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Artificial Sequence
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INFORMATION FOR SEQ ID NO:
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100.0%; Pred. No. 5.1e+02;
live 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: FastENG for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/038,637
FILING DATE: 10-MAR-1998
PRIOR APPLICATION NUMBER: 08/579,233
FILING DATE: 28-DEC-1995
APPLICATION NUMBER: 08/152,313
FILING DATE: 12-NOV-1993
ATTORNEY/AGENT INFORMATION:
NAME: Halle, Lisa A
REGISTRATION NUMBER: 38,347
REFERENCE/DOCKET NUMBER: 07265/146001
TELECOMMULICATION INFORMATION:
TELECOMMULICATION INFORMATION:
          CLASSIFICATION:
PRIOR APPLICATION:
PRIOR APPLICATION NUMBER: 08/107,673
FILING DATE: 18-Aug-93
APPLICATION NUMBER: 08/392,019
FILING DATE: 21-Feb-95
APPLICATION NUMBER: 08/571,047
FILING DATE: 12-Dec-95
APPLICATION NUMBER: 08/571,047
FILING DATE: 20-Dec-95
APPLICATION NUMBER: 08/55,191
FILING DATE: 22-Apr-1996
APPLICATION NUMBER: 08/635,820
FILING DATE: 22-Apr-1996
ATTORNEY/AGENT INFORMATION:
NAME: Friedman, Mark M.
REGISTRATION NUMBER: 33,883
REFERENCE/DOCKET NUMBER: 205/15
TELECOMMUNICATION INWERATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2122 CCCAGGCTGGAGTGCAG 2138
                                                                                                                                                                                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO: 2: SEQUENCE CHARACTERISTICS: LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity 100.
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: 972-3-5625554
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COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        619/678-5099
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; TOPOLOGY: linear
US-09-100-104-2
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US-08-837-201C-25
i Sequence 25, Application US/08837201C
j Fatent No. 598559
i GENERAL INFORMATION:
APPLICANT: Micholas M. Dean; Robert A. McKay; Loren J.
APPLICANT: Micholas M. Brenda F. Baker
i TITLE OF INVENTION: Compositions and Methods for the Modulation of
ITILE OF INVENTION: Compositions and Methods for the Modulation of
ITILE OF INVENTION: Activating Protein 1
NUMBER OF SEQUENCES: 139
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ö
                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 1; Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2; Indels
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COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MB STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 16.8; DB 1
Pred. No. 5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
0.7%; Score 16.8; D
Best Local Similarity 90.0%; Pred. No. 5e+0
Matches 18; Conservative 0; Mismatches
                                                                                       NAME: Altman, Daniel E
REGISTRATION NUMBER: 34,115
REPERENCE/DOCKET NUMBER: EIKENI.001APC
TELECOMINICATION INFORMATION:
TELEPHONE: 714-760-0404
TELEFAX: 714-760-9502
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM PS/2
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/837,201C
FILING DATE: APTil 14, 1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/JP95/02734
FILING DATE: 27-DEC-1995
ATTORNEY/AGENT INFORMATION:
NAME: Altman, Daniel E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2187 ATTCTCCTGCCTCAGCCTCC 2206
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ATTORNEY/AGENT INFORMATION:
NAME: Jane Massey Licata
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH.
TELECOMUNICATION INFORMATION:
TELEPHONE: (609) 810-11515
TELEFRAX: (609) 810-11515
TELEFRAX: (609) 810-1515
                                                                                                                                                                                                                            TELEX:
INFORMATION FOR SEQ ID NO: 1
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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TOPOLOGY: Linear
ANTI-SENSE: Yee
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US-08-849-701-12
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Sequence 12, Application US/09357073

Sequence 12, Application US/09357073

Sequence 12, Application US/09357073

Septicant: NFORMATION:

APPLICANT: Brett P. Monia

APPLICANT: Lex M. Cowsert

TITLE OF INVENTION: AND MAP KINASE KINASE 6 EXPRESSION

FILE REFERENCE: RTS-0086

CURRENT APPLICATION NUMBER: US/09/357,073
                                                          Gaps
                                                                                                                                                                                                                                                                                                JOS-09-01-05-55, Application US/09073567

Sequence 25, Application US/09073567

Patent No. 6013786

GENERAL INFORMATION:

APPLICANT: Jiandong Chen
APPLICANT: Sudhir Agrawal
APPLICANT: Sudhir Agrawal
APPLICANT: Ruiwen Zhang

TITLE OF INVENTION: MDM2-SPECIFIC ANTISENSE OLIGONUCLEOTIDES
NUMBER OF SEQUENCES: 49
CORRESPONDENCE ADDRESS:
ADDRESSEE: McDonnell Boehnen Hulbert & Berghoff
STREET: 300 South Wacker Drive, 32nd Floor
CITY: Chicago
STATE: IL
COUNTRY: United States of America
ZIP: 60606
COMPUTER: FRADABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: Microsoft Word 97
CURRENT APPLICATION NUMBER: US/09/073,567

FILING DATE:
CLASSIFICATION NUMBER: US/09/073,567

ATTORNEY/AGENT INFORMATION:
MATTORNEY/AGENT INFORMATION:
MATTORNEY/AGENT INFORMATION:
MATTORNEY/AGENT NUMBER: 98,057-A
TELECOMMUTCATION NUMBER: 98,057-A
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5e+02;
        DB 1; Length 20;
                                                          Indels
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Query Match 0.7%; Score 16.8; DB 1
Best Local Similarity 90.0%; Pred. No. 5e+02;
Matches 18; Conservative 0; Mismatches
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Best Local Similarity 90.0%; Pred. No. 5e+0
Matches 18; Conservative 0; Mismatches
                                                                                                                  2326 CCCACCTCGGCCTCCCAAAG 2345
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: nucleic acid HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (312) 913-0001
TELEPAX: (312) 913-0002
INFORMATION FOR SEQ ID NO: 25
SEQUENCE CHARACTERISTICS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: both
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
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ANTI-SENSE:

US-09-073-567-25
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Gaps
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                                                                                                     US-09-435-296-81/c

| Sequence B1, Application US/09435296
| Sequence B1, Application US/09435296
| GENERAL INFORMATION:
| APPLICANT: Brenda F. Baker
| APPLICANT: Brenda F. Baker
| TITLE OF INVENTION: ANTISENSE MODULATION OF RANK EXPRESSION
| FILE REPERENCE: RTS-0116
| CURRENT APPLICATION NUMBER: US/09/435,296
| CURRENT FILING DATE: 1999-11-05
| SEQ ID NO 81
| SEQ ID NO 81
| LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-280-805-26/c

; Sequence 26, Application US/09280805
; Patent No. 6184212
; Patent No. 6184212
; APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
; APPLICANT: Graham, Brett P. Monia
; TITLE OF INVENTION: EXPRESSION
; TITLE OF INVENTION: EXPRESSION
; ONDERSPONDENCE ADDRESS:
; CARRESPONDENCE ADDRESS:
; ADDRESSEE: Law Offices of Jane Massey Licata
; CITY: Marlton
; CITY: Marlton
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 16.8; DB 1;
Pred. No. 5e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         ) OTHER INFORMATION: Antisense Oligonucleotide US-09-435-296-81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0: Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: ISBN PC
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDFERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/280,805
FILING DATE: herewith
CLASSIFFCATION:
PRICK APPLICATION UNHBER: 09/048,810
FILING DATE: March 26,1998
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Licata, Jane Massey
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0346
TELECOMMUNICATION INFORMATION:
TELEPHONE: 609-810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2345 GIGCIGGGATTACAGGCAIG 2364
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                        20 CCAGCCTCGGCCTCCCAAAG 1
                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.7%;
Best Local Similarity 90.0%;
Matches 18; Conservative
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: Nucleic Acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETT
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Setent No. 613008

GRERAL INFORMATION:

GREEAL INFORMATION:

GREEAL INFORMATION:

TILLE OF INVENTION: ANTISENSE MODULATION OF TELOMERIC REPEAT BINDING FACTOR 1 EXPRESS

CURRENT APPLICATION NUMBER: US/09/358,384

CURRENT FILING DATE: 1999-07-21

SEQ ID NO 38

LENGTH: 20
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0.7%; Score 16.8; DB 1; Length 20;
Best Local Similarity 90.0%; Pred. No. 5e+02;
Matches 18; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                    Length 20;
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Patent No. 6171860
GENERAL INFORMATION:
APPLICANT: Brenda P. Baker
APPLICANT: Lex M. COWSET
TITLE OF INVENTION: ANTISENSE MODULATION OF RANK EXPRESSION
FILE REPRENCE: RTS-0116
CURRENT APPLICATION NUMBER: US/09/435,296
CURRENT APPLICATION DATE: 1999-11-05
NUMBER OF SEQ 1D NOS: 89
                                                                                                                                                                                                                                                             2; Indels
                                                                                                                                                                                                                  DB 1;
                                                                                                                                                                                                                  Score 16.8; DB :
Pred. No. 5e+02;
                                                                                                                                                   ; OTHER INFORMATION: Antisense Oligonucleotide US-09-357-073-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Antisense Oligonucleotide US-09-358-384-38
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ) OTHER INFORMATION: Antisense Oligonucleotide US-09-435-296-80
                                                                                                                                                                                                                Query Match
0.7%; Score 16.8; Di
Best Local Similarity 90.0%; Pred. No. 5e+0.
Matches 18; Conservative 0; Mismatches
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CURRENT FILING DATE: 1999-07-19
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ORGANISM: Artificial Sequence
                                                                                   TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA ORGANISM: Artificial Sequence
                  NUMBER OF SEQ ID NOS: 47
SEQ ID NO 12
LENGTH: 20
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-09-358-384-38/c
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LENGTH: 20
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GENERAL INFORMATION:
APPLICANT: Sidransky, David
TITLE OF INVENTION: DETECTION OF NEOPLASIM BY ANALYSIS OF SALIVA
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 4225 Executive Square, Suite 1400
CITY: La Jolla
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                                                                                           0.7%; Score 16.8; DB 1; Length 20; 90.0%; Pred. No. Se+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.7%; Score 16.8; DB.1; Length 20; 90.0%; Pred. No. 5e+02; ive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 26, Application US/09048810 Patent No. 6238921 GENERAL INFORMATION:
APPLICANT: Loren J. Miraglia, Pamela Nero, Mark J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IDEN Compatible
COMPUTER: IDEN Compatible
COMPUTER: SYSTEM: Windows 95
SOFTWARE: FASTERO FOR Windows Version 2.0b
CURRENT APPLICATION DATA:
FILING DATE: 10-MAR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       07265/146001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/579,233
FLING DATE: 28-DEC-1995
APPLICATION NUMBER: 08/152,313
FILING DATE: 12-NOV-1993
ATTORNEY/AGENT INFORMATION:
NAME: Haile, Lisa A.
REGISTRATION NUMBER: 38,347
                                                                                                                                                                     1695 TTTACATGTGCAAAGAAGCT 1714
                                                                                                                                                                                                                                                                                                      Sequence 135, Application US/09038637
Patent No. 6235470
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2110 CTTGCTCTGTTACCCAGGCT 2129
                                                                                                                                                                                                        20 TTTACATGTATAAGAAGCT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 07:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619/678-5070
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                                                                                         Query Match
Best Local Similarity 90.04
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 619/678-5099
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18; Conservative
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           single
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                     Linear
No
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Matches 18; Conserv
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USA
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                 ; TOPOLOGY:
; ANTI-SENSE:
US-09-280-805-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: La
STATE: CA
COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
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Gaps
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APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF CASPASE 8 EXPRESSION
FILTE OF INVENTION: ANTISENSE MODULATION OF CASPASE 9 CURESTERIOR
FILE REFERENCE: RTS-0107
CURRENT APPLICATION NUMBER: US/09/487,445
CURRENT FILING DATE: 2000-01-19
NUMBER OF SEQ ID NOS: 176
LENGTH: 20
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Pred. No. 5e+02;
0; Mismatches 2; Indels
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Brett P. Monia
ANTISENSE OLIGONUCLEOTIDE
MODULATION OF HUMAN MDM2 EXPRESSION
                                                                                                                                                                                                                                                                    DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
                                                                                   CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: WINDOWS FOR WORKGROUPS SOFTWARE: WORDPERFECT 5.1 CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/048,810 FILING DATE: herewith CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Antisense Oligonucleotide US-09-487-445-95
                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:

NAME: Licata, Jane Massey

REGISTRATION UNDRER: 32,237

REPERENCE/DOCKET NUMBER: ISPH-0302

TELECOMMUNICATION INFORMATION:

TELEPHONE: 609-810-1454

INFORMATION FOR SEQ ID NO: 26:

SEQUENCE CHARACTERISTICS:

LENGTH: 20 Dass pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1695 TITACATGTGCAAAGAAGCT 1714
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            2144 GATCTTGGCTCACTGCAAGC 2163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 95, Application US/09487445
Patent No. 6258600
GENERAL INFORMATION:
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Best Local Similarity 90.0°
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                             COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETT
                    TITLE OF INVENTION:
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                           COMPUTER: IBM 486
  Graham,
                                                                                                                                                                         STATE: NJ
COUNTRY: U.S.A.
ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TOPOLOGY: Linear
TI-SENSE: NO
                                                                                                                                  STREET: 66 Eag
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US-09-048-810-26
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APPLICANT: Brett P. Monia
APPLICANT: Brett P. Monia
APPLICANT: Robert McKay
APPLICANT: Robert McKay
APPLICANT: Madeline M. Butler
APPLICANT: Jacqueline Wyatt
TITLE OF INVENTION:
FILE REFERENCE: RTS-0115
CURRENT APPLICATION NUMBER: 105/9488,856A
CURRENT APPLICATION TO SEQ ID NOS: 88
NUMBER OF SEQ ID NOS: 88
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                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 MB STORAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.7%; Score 16.8; DB 1; 18est Local Similarity 90.0%; Pred. No. 5e+02; Matches, 18; Conservative 0; Mismatches 2;
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90.0%; Pred. No. 5e+02;
tive 0; Mismatches 2;
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                                                                                                                                                                                                                                                                  CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/837,201
FILING DATE: APPLI 14, 1997
ATTORNEY/AGENT INFORMATION:
NAME: Jane Massey Licata
REGISTATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0209
TELECOMMUNICATION INFORMATION:
                                                                                                                             COMPUTER: IBM PS/2
OPERATING SYSTEM: WINDOWS 95
SOFTWARE: WORDPERFECT 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/364,416
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 73, Application US/09488856A; Patent No. 6316259; GENERAL INFORMATION:
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (609) 810-1515
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (609) 810-1454
INFORMATION FOR SEQ ID NO: 2
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 90.0
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Linear
Marlton
                                                                                                                                                                                                                                                      FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-488-856A-73/c
                                                                ZIP: 08053
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; ANTI-SENSE:
US-09-364-416-25
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LENGTH: 20
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                                                 Sequence 70. Application US/09467642

Patent No. 6300132

GENERAL INFORMATION:

APPLICANT: Bratt P. Monia

APPLICANT: Lex M. Cowsert

TITLE OF INVENTION: ANTISENSE MODULATION OF TELOMERIC REPEAT BINDING FACTOR 2 EXPRES

FILE REFERENCE: RTS-0106

CURRENT FILING DATE: 1999-12-20

NUMBER OF SEQ ID NOS: 89
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APPLICANT: Brett P. Monia

APPLICANT: Lex M. COMBERT

TITLE OF INVENTION: ANTISENSE MODULATION OF TELOMERIC REPEAT BINDING FACTOR 2 EXPRES

FILE REPERENCE: RTS-0106

CURRENT APPLICATION NUMBER: US/09/467,642

CURRENT FILING DATE: 1999-12-20

NUMBER OF SEQ ID NOS: 89

SEQ ID NO 72

LENGTH: 20
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Retent No. 6312900

GENERAL INFORMATION:
APPLICANT: Nicholas M. Dean; Robert A. McKay; Loren J. APPLICANT: Miraglia; Brenda F. Baker
TITLE OF INVENTION: Antisense Oligonucleotide
TITLE OF INVENTION: Compositions and Methods for the Modulation of
TITLE OF INVENTION: Activating Protein 1
NUMBER OF SEQUENCES: 139
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0.7%; Score 16.8; DB 1; Length 20; illarity 90.0%; Pred. No. 5e+02; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                               ; FEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-09-467-642-70
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Antisense Oligonucleotide US-09-467-642-72
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Patent No. 6300132
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                                                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Artificial Sequence
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Best Local Similarity 90.0
Matches 18; Conservative
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Best Local Similarity
Matches 18; Conserv
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                                        -09-467-642-70/c
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                                                                                                                                                                                                                                                                                        SEQ ID NO 70
LENGTH: 20
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APPLICANT: SIMON, JASON
APPLICANT: SIMON, JASON
APPLICANT: BANDIN, JASON
APPLICANT: BANDIN, SUNIL
TITLE CANT: PANDIN, SUNIL
TITLE OF INVENTION: NOVEL HUMAN GENES RELATING TO RESPIRATORY DISEASES AND OBESITY
FILE REFERENCE: 2976-4037
CURRENT APPLICATION NUMBER: 06/211,749
PRIOR FILING DATE: 2000-06-14
PRIOR FILING DATE: 1999-07-30
NUMBER OF SEQ ID NOS: 51
SOFTWARE: PATENTI NOT 2.1
SEQ ID NO 2.7
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Vickie L. Brown-Driver
APPLICANT: Vickie L. Brown-Driver
APPLICANT: Hong Zhang
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF CASPASE 6 EXPRESSION
FILLE REPERENCE: RTS-0187
CURRENT APPLICATION NUMBER: US/09/679,299A
CURRENT FILING DATE: 2000-10-04
NUMBER OF SEQ ID NOS: 164
LENGTH: 20
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                                                                                                                                                                           Query Match 0.7%; Score 16.8; DB 1; Length 20; Best Local Similarity 90.0%; Pred. No. 5e+02; Matches 18; Conservative 0; Mismatches 2; Indels
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                                                                                                      ), OTHER INFORMATION: Antisense Oligonucleotide US-09-657-346A-24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ) OTHER INFORMATION: Antisense Oligonucleotide US-09-679-299A-71
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-627-465B-27
; Sequence 27, Application US/09627465B
; Patent No. 6737519
                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 71, Application US/09679299A Patent No. 6566135
                                                                                                                                                                                                                                                                               2193 CTGCCTCAGCCTCCCAATTA 2212
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APPLICANT: KEITH, TIM
APPLICANT: LITTLE, RANDALL D.
APPLICANT: EERDEWEGH, PAUL VAN
APPLICANT: DUPUIS, JOSEE
APPLICANT: DEL MASTRO, RICHARD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Artificial Sequence
                                  TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Artificial Sequence
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            LENGTH: 20
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                                                                                    FEATURE:
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Patent No. 6410325

GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: Sugan M. Freier
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: EXPRESSION
FILE REPRENEUR: RY. 2020
CURRENT APPLICATION NUMBER: US/09/851,896
CURRENT FILING DATE: 2001-05-08
NUMBER OF SEQ ID NOS: 89
SEQ ID NOS: 89
US-09-662-250A-76

Sequence 76, Application US/09662250A

Sequence 76, Application US/09662250A

Sequence 76, Application US/09662250A

GENERAL INFORMATION:

APPLICANT: Breet P. Monia

APPLICANT: Jacqueline Wyatt

APPLICANT: APPLICANT: ANTISENSE MODULATION OF PHOSPHORYLASE KINASE BETA EXPRESSION

TITLE OF INVENTION: ANTISENSE MODULATION OF PHOSPHORYLASE KINASE BETA EXPRESSION

FILE REFERENCE: RTS-0129

CURRENT APPLICATION NUMBER: US/09/662,250A

CURRENT FILING DATE: 2000-09-14

NUMBER OF SEQ ID NOS: 102

SEQ ID NO 76

LENGTH: 20
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Sequence 24, Application US/09657346A
Patent No. 6503754
GENERAL INFORMATION:
APPLICANT: Hong Zhang
APPLICANT: Hong Zhang
APPLICANT: Hong Zhang
APPLICANT: Argueline Wyatt
TITLE OF INVENTION: EXPRESSION
FILE REFERENCE: RTS-0135
CURRENT APPLICATION NUMBER: US/09/657,346A
SEQ ID NOS: 174
SEQ ID NOS: 174
SEQ ID NOS: 174
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2; Indels
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Best Local Similarity 90.0%; Pred. No. 5e+02;
Matches 18; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FEATURE:

COTHER INFORMATION: Antisense Oligonucleotide
US-09-851-896-17
                                                                                                                                                                                                                                                                                                                                                                                          ) OTHER INFORMATION: Antisense Oligonucleotide US-09-662-250A-76
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2115 TCTGTTACCCAGGCTGGAGT 2134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2344 AGTGCTGGGATTACAGGCAT 2363
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 90.04
Matches 18; Conservative
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US-09-657-346A-24/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 570
US-09-851-896-17/c
                                                                                                                                                                                                                                                                                                                         TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
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FITLE OF INVENTION: REPEAT DNA MARKERS
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                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Schum, James W.
APPLICANT: Schum, Jeffery W.
TITLE OF INVENTION: MATERIALS AND METHODS FOR
TITLE OF INVENTION: IDENTIFYING AND ANALYZING INTERMEDIATE TANDEM
TITLE OF INVENTION: REPEAT DNA MARKERS
NUMBER OF SEQUENCES: 147
CORRESPONDENCE ADDRESS:
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MATERIALS AND METHODS FOR
IDENTIFYING AND ANALYZING INTERMEDIATE TANDEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ö
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                     ; OTHER INFORMATION: Description of Artificial Sequence: Primer US-09-627-465B-27
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                                                                                      Query Match 0.7%; Score 16.8; DB 1; Length 20; Best Local Similarity 90.0%; Pred. No. 5e+02; Matches 18; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: Wisconsin
COUNTRY: U.S.A.
ZIP: 53711-5399
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb
COMPUTER: IBM compatible PC
OPERATING SYSTEM: Windows 95
SOFTWARE: Word 97 (DOS text format)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/018,584A
FILING DATE: 04-Feb-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY AGENT INFORMATION:
NAME: Grady J. Frenchick
REGISTRATION NUMBER: 29,018
REFERENCE/DOCKET NUMBER: 16026.9180
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 257-3501
TELEPHONE: (608) 257-2275
INFORMATION FOR SEQ ID NO: 67:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 112, Application US/09018584A Patent No. 623863
GENERAL INFORMATION:
APPLICANT: Bacher, Jeffery W. TITLE OF INVENTION: MATERIALS AND PITTLE OF INVENTION: INDENTIFYING ANI
                                                                                                                                                                                                                                                                                                                US-09-018-584A-67/c
; Sequence 67, Application US/09018584A
Patent No. 6238863
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2111 TIGCICIGITACCCAGGCIG 2130
                                                                                                                                                                               2347 GCTGGGATTACAGGCATGAG 2366
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Promega Corporation
STREET: 2800 Woods Hollow Road
                                                                                                                                                                                                                         1 GTTGGGATTACAGGCACGAG 20
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Best Local Similarity 90.0°
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION:
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STATE:
FEATURE:
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APPLICANT: Schumm, James W.
Bacher, Jeffery W.
TITLE OF INVENTION: MATERIES AND METHODS FOR REPEAT FANDEM REPEAT DNA MARKERS
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Madison
STATE: Wisconsin
COUNTY: U.S.A.
ZIP: 53711-5399
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb
COMPUTER: IBM compatible PC
COMPUTER: Word 97 (DOS text format)
CURRENT APPLICATION DATA:
                                                                                                                          CUNTRY: WISCALLER WINDOWS 95
COMPUTER: IBM compatible PC
OPERATING SYSTEM: Windows 95
SOFTWARE: Word 97 (DOS text format)
CURRENT APPLICATION DATA:
APPLICATION UNDER: US/09/018,584A
FILING DATE: 04-Peb-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/09/784,423
FILING DATE: 15-Feb-2001
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY AGENT INFORMATION:
NAME: Grady J. Frenchick
REGISTRATION NUMBER: 29,018
REFERENCE/DOCKET NUMBER: 16026.9180
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 257-251
TILEPHONE: (608) 257-2275
INFORMATION FOR SEQ ID NO: 112:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/018,584
FILING DATE: 04-Feb-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Promega Corporation
STREET: 2800 Woods Hollow Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2106 GAGTCTTGCTCTGTTACCCA 2125
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Patent No. 6767703
GENERAL INFORMATION:
                                           3: Promega Corporation 2800 Woods Hollow Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 20 GAGTCTTACTCTGTTGCCCA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 14
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: Nucleic Acid
STRANDEDNESS: Single
NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: Promega C
                                                                                                              STATE: Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Linear
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US-09-018-584A-112
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Sequence 11
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.7%; Score 16.8; DB 1; Length 21; 90.0%; Pred. No. 4.9e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0.7%; Score 16.8; DB 1; Length 21; 90.0%; Pred. No. 4.9e+02; Live 0; Mismatches 2; Indels
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CORRESONDENCE ADDRESS: 147
CORRESSED Promega Corporation
STREET: 2800 Woods Hollow Road
CITY: Madison
STATE: 4800 Woods Hollow Road
CITY: Madison
STATE: Wisconsin
COUNTRY: U.S.A.
ZIP: 53111-5399
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb
COMPUTER: IBM compatible PC
OMPUTER: IBM compatible PC
OMPUTER: IBM compatible PC
OMPUTER: Mord 97 (DOS text format)
COMPUTER: Word 97 (DOS text format)
APPLICATION DATA:
APPLICATION NUMBER: US/09/784,423
FILING DATE: 15-Feb-2001
CLASSIFICATION NUMBER: 09/018,584
FILING DATE: 04-Feb-1998
ATTORNEY/AGRAT INFORMATION:
NAMME: GYADA J. Frenchick
REGISTRATION NUMBER: 29,018
REFERENCE/DOCKET NUMBER: 16026.9180
                                 REGISTRATION NUMBER: 29,018
REFERENCE/DOCKET NUMBER: 16026.9180
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 257-3501
TELEFAX: (608) 257-2275
INFORMATION FOR SEQ ID NO: 67
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE DESCRIPTION: SEQ ID NO: 112
                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
SEQUENCE DESCRIPTION: 570 ID NO: 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2111 TIGCICIGITACCCAGGCTG 2130
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TELEPHONE: (608) 257-3501
NAME: Grady J. Frenchick
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: (608) 257-2275
INFORMATION FOR SEQ ID NO: 112
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   20 TIGCTCTGTCACCAAGGCTG 1
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STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 147
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 90.0 Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 90.0
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                             LENGTH: 21
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US-09-784-423-112/c
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                                                                                                                                                         US-08-117-952-287/c

| Sequence 287, Application US/08117952
| Patent No. 5851760
| GENERAL INFORMATION:
| APPLICANT: Evans, Glen A. APPLICANT: Brith, Michael W. TITLE OF INVENTION: METHOD FOR GENERATION OF SEQUENCE; TITLE OF INVENTION: SAMPLED MAPS OF COMPLEX GENOMES NUMBER OF SEQUENCES: 797
| CORRESPONDENCE ADDRESS: ADDRESS: STREET: 444 South Flower Street, Suite 2000 CITY: Los Angeles
| CITY: Los Angeles STATE: CA. COMPLEX CALLED COMPLEX GENOMES STREET: CA. COMPLEX GENOMES GENOM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER FRADABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: EN PC Compatible
COMPUTER: EN PC Compatible
COMPUTER: EN PC Compatible
CORERATIOG SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/117,952
FILING DATE: 07-SEP-1993
CLASSIFICATION NUMBER: US 08/078,471
PFILING DATE: 15-JUN-1993
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 31,192
REGISTRATION NUMBER: 941 9423
TELEPHONE: 619-546-4737
TELEFRAK: 619-546-4737
INFORMATION FOR SEQ ID NO: 287:
SEQUENCE CHARACTERISTICS:
LENGTH: 22 base pairs
TYPE: NUCLeic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 25, Application US/08859998
Patent No. 594076
GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
APPLICANT: Jokhadze, George
APPLICANT: Bibliashvilli, Robert
ITILE OF INVENTION: BETHOD OF ASSAXING DIFFERENTIAL
ITILE OF INVENTION: EXPRESSION
NUMBER OF SEQUENCES: 1375
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              20 GAGTCTTACTCTGTTGCCCA 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        USA
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US-09-225-928-25/c
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| Sequence 92, Application US/09245041 |
| Sequence 92, Application US/09245041 |
| Patent No. 6274339 |
| GENERAL INFORMATION: |
| APPLICANT: Moore, K. |
| APPLICANT: Moore, K. |
| APPLICANT: Moore, K. |
| TITLE OF INVENTION: OF BODY WEIGHT DISORDERS INCLUDING OBESITY |
| TITLE OF INVENTION: OF BODY WEIGHT DISORDERS INCLUDING OBESITY |
| TITLE OF INVENTION: OF BODY WEIGHT DISORDERS INCLUDING OBESITY |
| TITLE OF INVENTION: OF BODY WEIGHT DISORDERS INCLUDING OBESITY |
| CURRENT APPLICATION NUMBER: US/09/245,041 |
| CURRENT FILING DATE: 1999-02-05 |
| EARLIER PILING DATE: 1999-07-21 |
| EARLIER PILING DATE: 1999-10-20 |
| NUMBER OF SEQ ID NOS: 131 |
| SOSTWARE FEASEQ for Windows Version 3.0 |
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 22;
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0.7%; Score 16.8; DB 1;
Best Local Similarity 90.0%; Pred. No. 4.9e+02;
Matches 18; Conservative 0; Mismatches 2;
                                                                                                                                                                                   MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FRASESQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/859,998
FILING DATE: 21-MAY-1997
CLASSIFICATION: 435
PRIOR APPLICATION NUMBER: STANAY-1697
ATTORNEY, AGENT INFORMATION:
ARBITANAY GART INFORMATION:
NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 90906/002001
TELEPHONE: 415-322-5070
TELEPHONE: 415-322-5070
TELEPHONE: 415-322-5070
TELEPHONE: GIARACTERISTICS:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION: oligonucleotide primer
                    ADDRESSEE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2122 CCCAGGCTGGAGTGCAGTGG 2141
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; ORGANISM: Artificial Sequence
US-09-245-041-92
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 22 base pairs
                                     STREET: 2200 Sand Hill
CITY: Menlo Park
STATE: CA
COUNTRY: US
ZIP: 94025
COMPUTER READBALE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM Compatil
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ropology:
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LENGTH: 22
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; Sequence 25, Application US/09225201B
; Sequence 10.46489455
; Patent No. 6489455
; GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
JORAGLe, George
; Bibliashvilli, Robert
; TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
                                                                                                                                                   Sequence 25, Application US/09225928
Patent No. 6352829
GENERAL INFORMATION:
APPLICANT: Chenchik, Alex
Jokhadze, George
Bibliashvilli, Robert
TITLE OF INVENTION: METHOD OF ASSAYING DIFFERENTIAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM Compatible
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FEAGESC for Windows Version 2.0
SOFTWARE: FEAGESC for Windows Version 2.0
CURRENT APPLICATION NUMBER: US/09/225,928
FILING DATE: 05-Jan-1999
CLASSIFICATION NUMBER: 08/859,998
APPLICATION NUMBER: 08/859,998
ATTORNEY/AGENT: INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; SEQUENCE DESCRIPTION: oligonucleotide primer; SEQUENCE DESCRIPTION: SEQ ID NO: 25: US-09-225-928-25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.7%; Score 16.8; DB 1;
90.0%; Pred. No. 4.9e+02;
tive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                         NUMBER OF SEQUENCES: 1375
CORRESPONDENCE ADDRESS:
ADDRESSE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
CITY: Menlo Park
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 09096/002001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-322-5070
TELEPHONE: 415-322-5070
SEQUENCE CHARACTERISTICS:
2268 GAGACAGGGTTTCACCGTGT 2287
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2122 CCCAGGCTGGAGTGCAGTGG 2141
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 22 base pairs
                                          21 cacacaccicrcacrcrcrc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity 90.0
Matches 18; Conservative
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2268 GAGACAGGGTTTCACCGTGT 2287
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OTHER INFORMATION: Primer
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-893-238-92
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US-09-156-253-45
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APPLICANT: MOOFE, K.
APPLICANT: MOSE, K.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR THE DIAGNOSIS AND
TITLE OF INVENTION: TREATMENT OF BODY WEIGHT DISORDERS INCLUDING:
TITLE OF INVENTION: OBESITY
FILE REFERENCE: 7963-151
CURRENT APPLICATION NUMBER: US/09/358,055B
CURRENT APPLICATION NUMBER: 09/245,041
PRIOR APPLICATION NUMBER: 09/245,041
PRIOR RILING DATE: 1999-07-05
NUMBER OF SEQ ID NOS: 153
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 93
LENGTH: 22
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                                                                                                                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows95
SOFTWARE: FastSEQ for Windows Version 2.0
NUMBER OF SEQUENCES: 1375
CORRESPONDENCES: 1375
CORRESPONDENCES: 1375
CORPUTE: ADDRESSE: Fish & Richardson, P.C.
STREET: 2200 Sand Hill Road, Suite 100
CITY: Menlo Park
STATE: CA
COUNTRY: US
ZIP: 94025
COMPUTER READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/09/225,201B
FILING DATE: 05-Jan-1999
CLASSIFICATION: «Unknown»
PRIOR APPLICATION: «Unknown»
PRIOR APPLICATION: «Unknown»
APPLICATION NUMBER: US/08/859,998
FILING DATE: 21-MAY-1997
ATTORNEY/AGENT INFORMATION:
NAME: Field, Bret E.
REGISTRATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 39,6206/002001
TELECOMMUNICATION NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 37,620
REFERENCE/DOCKET NUMBER: 37,620
REFERENCE/DOCKET AUMBER: 37,620
RE
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US-09-358-055B-93/c
Sequence 93, Application US/09358055B
; Patent No. 6713277
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH: 22 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FEATURE:

OTHER INFORMATION: Primer
US-09-358-0558-93
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE
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Sequence 92, Application US/09893238

Sequence 92, Application US/09893238

Patent No. 6727348

GRERAL INFORMATION:
APPLICANT: Moore, K.
TITLE OF INVENTION:
TITLE OF INVENTION:
CURRENT APPLICATION NUMBER: US/09/893,238

CURRENT PILING DATE: 2010-06-27

PRIOR APPLICATION NUMBER: 09/245,041

PRIOR APPLICATION NUMBER: 09/245,041

PRIOR APPLICATION NUMBER: 60/245

PRIOR APPLICATION NUMBER: 60/093,630

PRIOR FILING DATE: 1998-07-21

PRIOR PRILING DATE: 1998-07-21

PRIOR PRILING DATE: 1998-10-20

NUMBER OF SEQ ID NOS: 129

SEQ ID NO 92

LENGTH: 22

LENGTH: 22
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                                                            Gaps
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US-09-156-253-45
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 45, Application US/09156253C
| Patent NO. 6001652
| GENERAL INFORMATION:
| APPLICANT: Baker, Breath P. |
| APPLICANT: Cowsert, Lex M. |
| APPLICANT: Cowsert, Lex M. |
| TITLE OF INVENTION: Antisense Modulation of CREL Expression FILE REFERENCE: RTS-0010 |
| CURRENT APPLICATION NUMBER: US/09/156,253C |
| CURRENT FILING DATE: 1998-09-18 |
| NUMBER OF SEQ ID NOS: 48 |
| SEQ ID NO 45 |
| LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.7%; Score 16.8; DB 1; Length 22; Best Local Similarity 90.0%; Pred. No. 4.9e+02; Matches 18; Conservative 0; Mismatches 2; Indels
Query Match 0.7%; Score 16.8; DB 1; Length 22; Best Local Similarity 90.0%; Pred. No. 4.9e+02; Matches 18; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 18;
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Best Local Similarity 94.4%; Pred. No. 5.4e+02;
Matches 17; Conservative 0; Mismatches 1;
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19 ATTACAGACATGAGCCAC
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TITLE OF INVENTION: id.
TITLE OF INVENTION: su.
NUMBER OF SEQUENCES: 2:
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: DC
ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                          CITY: Raleigh
STATE: NC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-07-952-442-19/c
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APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Matches
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US-06-30-706-94

Sequence 94, Application US/09630706

Patent No. 6277640

GENERAL INFORMATION:

APPLICANT: C. Frank Bennett

APPLICANT: C. Frank Bennett

TITLE OF INVENTION: ANTISENSE MODULATION OF HER-3 EXPRESSION

FILE REFERENCE: RTS-0053

CURRENT APPLICATION WHERE: US/09/630,706

CURRENT FILING DATE: 2000-08-01

NUMBER OF SEQ ID NOS: 94

LENGTH: 18
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                                                                                                                                Sequence 47, Application US/09161443A
Patent No. 6020198
GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: Lex M. COWSERT
TITLE OF INVENTION: ANTISENSE MODULATION OF RIP-1 EXPRESSION
FILE REFERENCE: RTS-0011
CURRENT APPLICATION NUMBER: US/09/161,443A
CURRENT FILING DATE: 1998-09-25
NUMBER OF SEQ ID NOS: 47
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.7%; Score 16.4; DB 1; Length 18; Best Local Similarity 94.4%; Pred. No. 5.4e+02; Matches 17; Conservative 0; Mismatches 1; Indels
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Patent No. 6045997
GENERAL INFORMATION:
APPLICANT: Futreal, Phillip A
APPLICANT: Ashworth, Alan
APPLICANT: Statton, Michael R
TITLE OF INVENTION: Materials and methods relating to the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1; Indels
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0.7%; Score 16.4; DB 1;
Best Local Similarity 94.4%; Pred. No. 5.4e+02;
Matches 17; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: Antisense Oligonucleotide US-09-161-443-47
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Antisense Oligonucleotide US-09-630-706-94
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2336 CCTCCCAAAGTGCTGGGA 2353
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2341 CAAAGTGCTGGGATTACA 2358
  2338 TCCCAAAGTGCTGGGATT 2355
                       1 TCCCAAAGTGCTAGGATT 18
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                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             RESULT 588
US-08-755-587-124/c
                                                                                                 RESULT 586
US-09-161-443-47/c
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identification and sequencing of the BRCA2 cancer susceptibility gene and uses thereof.
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                                                                    ADDRESSEE: Bell Seltzer Park & Gibson
STREET: 310 UCB Plaza, 3605 Glenwood Avenue, PO Drawer 31107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Soubrier, Florent
APPLICANT: Cotelevtsev, Youri
APPLICANT: Corval, Pierre
ITILE OF INVENTION: Angiotensinogen Gene Variants and
TITLE OF INVENTION: Predisposition to Essential Hypertension
NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 19;
                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ADDRESSEE: Venable, Baetjer, Howard & Civiletti
STREET: 1201 New York Avenue N.W., Suite 1000
CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0.7%; Score 16.4; DB 1;
94.4%; Pred. No. 5.3e+02;
tive 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 31,665
REFERENCE/DOCKET NUMBER: 5405-135
INFORMATION FOR SEQ ID NO: 124:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
TYPE: mucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 19, Application US/07952442; Patent No. 5374525; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2353 ATTACAGGCATGAGCCAC 2370
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Jeunemaitre, Xavier
Lifton, Richard P.
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APPLICANT: Hiroteu, Takuo
APPLICANT: Hiroteu, Takuo
APPLICANT: Keshi, Hiroyuki
APPLICANT: Matsuhisa, Akio
TITLE OF INVENTION: Probe for Diagnosing Infectious Diseases
NUMBER OF SEQUENCES: 8
CORRESPONDENCE ADDRESS:
ADDRESSEE: Marshall, O'Toole, Gerstein, Murray & Borun
STREET: 6300 Sears Tower, 233 South Wacker Drive
                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 0.7%; Score 16.4; DB 1; Length 20; Best Local Similarity 94.4%; Pred. No. 5.3e+02; Matches 17; Conservative 0; Mismatches 1; Indels
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MEDIUM TYPE: FODDY disk
COMPUTER READBLE FORM:
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.30
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/416,831B
FILING DATE: 19-UW-1995
PRIOR APPLICATION NUMBER: 25-CCT-1993
PRIOR APPLICATION NUMBER: 35-CCT-1993
PRIOR APPLICATION NUMBER: 38,153
FILING DATE: 23-CCT-1993
ATTORNEY/AGENT INFORMATION:
NAME: Gass, David A.
RESISTANTION NUMBER: 19036/32578
TELEFROM UNIMBER: 38,153
REFERENCE/DOCKET NUMBER: 19036/32578
TELEFROM UNIMBER: 312/474-6340
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "Synthetic DNA"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET: 630v CITY: Chicago STATE: 1111nois COUNTRY: United States of America ZIP: 60605-6402 ZIP: GANDABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 5, Application US/08416831B Patent No. 5708159 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2122 CCCAGGCTGGAGTGCAGT 2139
                                                                                                         TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
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                                                                                                                                                                                                                                                                                                                                 Homo sapiens
    LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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Best Local Similarity
                                                                                                                                                                                                       HYPOTHETICAL: N
ANTI-SENSE: NO
ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         JS-08-416-831B-5/c
                                                                                                                                                                                                                                                                                                                     ) ORGANISM:
US-08-269-766-19
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APPLICANT: Jeunemaitre, Xavier
APPLICANT: Jeunemaitre, Xavier
APPLICANT: Liften, Richard P.
APPLICANT: Liften, Richard P.
APPLICANT: Kotelevesev, Youri
APPLICANT: Kotelevesev, Youri
APPLICANT: Corval, Pierre
TITLE OF INVENTION: Angiotensinogen Gene Variants and
TITLE OF INVENTION: Angiotensinogen Gene Variants and
TITLE OF INVENTION: Predisposition to Essential Hypertension
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
ADDRESSEE: Venable, Baetjer, Howard & Civiletti
STREET: 1201 New York Avenue N.W., Suite 1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.7%; Score 16.4; DB 1; Length 20; 94.4%; Pred. No. 5.3e+02; tive 0; Mismatches 1; Indel8
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STATE: DC
ZIP: 2006
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: TEM PC compatible
COMPUTER: TEM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/269,766
FILING DATE: 01-JUL-1994
CLASSIPICATION: DATA:
APPLICATION NUMBER: US 07/952,442
FILING DATE: 30-SEP-1992
FILING DATE: 30-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAME: INDEM, Jeffrey L.
REGISTRATION NUMBER: 28,957
                                                                                                                                                                                                                                            19780-104502
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APPLICATION NUMBER: US/07/952,442 FILING DATE: 19920930 CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 19, Application US/08269766
Patent No. 5589584
                                                                 CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Inhen, Jeffrey L.
REGISTRATION NUMBER: 28,957
REFERENCE/DOCKET NUMBER: 19780
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION:
TELEC
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TELECOMMUNICATION INFORMATION:
TELEX: 202-962-4810
TELEX: 202-962-8300
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          18 CCGAGGCTGGAGTGCAGT 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 94.4 Matches 17; Conservative
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US-08-269-766-19/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
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Jean-Marc

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APPLICANT: Kotelevteev, Youri
APPLICANT: Corvol, Pierre
TITLE OF INVENTION: Method to Determine Predisposition
TITLE OF INVENTION: to Hypertension
NUMBER OF SEQUENCES: 22
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Venable, Baetjer, Howard & Civiletti
STREET: 1201 New York Avenue N.W., Suite 1000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: DC.

ZIP: 20005

COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 5.1/5.2 Windows
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/092,988
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFCATION:
CLASSIFCATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/319,545
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Ihnen, Jeffrey L.
REGISTRATION NUMBER: 28,957
REPERENCE/DOCKET NUMBER: 19780-1
TELECOMMUNICATION INFORMATION:
TELECHONE: 202-962-4810
TELECT: 202-962-4810
                                                         Jeunemaitre, Xavier
Lifton, Richard P.
                                                                                                                                              Soubrier, Florent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STREET: 1201 New CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ANTI-SENSE: NO ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               HYPOTHETICAL:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        US-09-092-988-19
                                                                                               APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 594
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    1; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Jeunemaitre, Xavier
APPLICANT: Lifton, Richard P.
APPLICANT: Soubrier, Florent
APPLICANT: Scolevtsev, Youri
APPLICANT: Corvol, Pierre
APPLICANT: Corvol
APPLICANT: 
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Venable, Baetjer, Howard & Civiletti
STREET: 1201 New York Avenue N.W., Suite 1000
CITY: Washington
STATE: DC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STATE: DC
ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 5.1/5.2 Windows
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/319,545A
FILING-DATE: 7-OCT-1994
CLASSIFICATION: 435
FILING-DATE: 30-SEP-1992
ATTORNEY, AGENT INFORMATION:
NAME: INDEM, Jeffrey L.
REGISTRATION NUMBER: 28,957
REFERENCE/DOCKET NUMBER: 19780-104502-2
REGISTRATION NUMBER: 28,957
REFERENCE/DOCKET NUMBER: 19780-104502-2
TELECOMUTINICATION INFORMATION:
TELEPHONE: 202-962-4810
FIELERHONE: 202-962-8310
FIELERHONE: 202-962-8310
FIELERHONE: 202-962-8310
FIELERHONE: 202-962-8310
FIELERHONE: 202-962-8310
FIELERHONE: 302-962-8310
FIE
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    0; Mismatches
                                                                                                                                                                                                                                                                                                                                                           US-08-319-545A-19/c
; Sequence 19, Application US/08319545A
; Patent No. 2763168
; GENERAL INFORMATION:
APPLICANT: Lalouel, Jean-Marc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-092-988-19/c
; Sequence 19, Application US/09092988
Patent No. 5998145
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2122 CCCAGGCTGGAGTGCAGT 2139
                                                                                          393 GTTAGACCAAAGCCATTG 410
                                                                                                                                              20 GTTAGACCTAAGCCATTG 3
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MOLECULE TYPE: DNA (genomic)
HYPOTHETICAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.7%;
Best Local Similarity 94.4%;
Matches 17; Conservative
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ORIGINAL SOURCE:
ORGANISM: Homo sapiens
17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-319-545A-19
Matches
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19780-104502-2

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Pred. No. 5.3e+02;
0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 19, Application US/09106216
Patent No. 6153386
GENERAL INFORMATION:
APPLICANT: Lalouel, Jean-Marc
APPLICANT: Jennemaire, Xavier
APPLICANT: John Richard P.
APPLICANT: Soubrier, Florent
APPLICANT: Rotelevrsev, Yuri
APPLICANT: Rotelevrsev, Yuri
TITLE OF INVENTION: Method to Determine Predisposition to NUMBER OF ENVENTION: Hypertension
MANDED OF SEVENTION: Hypertension
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                E: Rothwell, Figg, Ernst & Kurz
555 Thirteenth Street N.W., Suite 701-E
                                                                                                                                                                                                                                                                                                                                                                 2122 CCCAGGCTGGAGTGCAGT 2139
                                                                            TOPOLOGY: linear MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                 Query Match 0.7%;
Best Local Similarity 94.4%;
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                              18 ccgaggcrggagrgcagr
                                                                                                                                                                                                   Homo sapiens
LENGTH: 20 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: CORRESPONDENCE ADDRESS: ADDRESSE: Rothwell,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 77, Application US/10215448
Patent No. 6716975
GENERAL INFORMATION:
APPLICAMT: GACQUELINE Wyatt
TITLE OF INVENTION: ANTISENSE MODULATION OF EDG1 EXPRESSION
FILE REPREMENTE: RTS-0179
CURRENT APPLICATION NUMBER: US/10/215,448
CURRENT FILING DATE: 2002-08-09
NUMBER OF SEQ ID NOS: 105
EBNGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score 16.4; DB 1;
Pred. No. 5.3e+02;
0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; FEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-10-215-448-77
                                                                                                                                                                                                                                                                                             19780-104502-2
                                                                 PRIOR APPLICATION:
PRIOR APPLICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 7-0CT-1994
APPLICATION NUMBER: US 07/952,442
FILING DATE: 30-SEP-1992
ATTORNEY/AGENT INFORMATION:
NAMB: Inhen, Jeffrey L.
REGISTRATION NUMBER: 28,957
REGISTRANICH/DOKET NUMBER: 19780-10450:
TELEPHONE: 202-962-4810
TELEPHONE: 202-962-4810
TELEPHONE: 202-962-4810
TELEPHONE: 202-962-8300
INFORMATION FOR SEQ ID NO: 19:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/429,034
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 6, Application US/08394210; Patent No. 5814716; GENERAL INFORMATION:
APPLICANT: JALLAT, SOPHIE
APPLICANT: MEULIEN, PIERRE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2122 CCCAGGCTGGAGTGCAGT 2139
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
HYPOTHFILAL: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match 0.7%;
Best Local Similarity 94.4%;
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                   LENGTH: 20 base paire
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ANTI-SENSE: NO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ORIGINAL SOURCE;
ORGANISM: Hor
US-09-429-034-19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-10-215-448-77/C
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Jounealite, Xavier
APPLICANT: Jounealite, Xavier
APPLICANT: Jounealite, Xavier
APPLICANT: Jounealite, Xavier
APPLICANT: Jounealite, Richard P.
APPLICANT: Jounealite, Florent
APPLICANT: Kotelevtsev, Youri
APPLICANT: Kotelevtsev, Youri
APPLICANT: Corvol, Plaire
TITLE OF INVENTION: Method to Determine Predisposition
TITLE OF INVENTION: to Hypertension
NUMBER OF SEQUENCES: 22
CORRESPONDENCES: 22
CORRESPONDENCES: 32
ADDRESSE: Venable, Baetjer, Howard & Civiletti
STREE: 1201 New York Avenue N.W., Suite 1000
CITY: Weahington
STATE: DC
ZIP: 20005
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM FC Compatible
COMPUTER: IBM FC Compatible
COMPUTER: IBM FC COMPATISH: PC-DOS/MS-DOS
SOFTWARE: WordPerfect 5.1/5.2 Windows
                                                                    ZIP: 2004
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/106,216
FILING DATE:
                                                                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 09/
FILING DATE: 08-UN-1998
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/319,545
FILING DATE: 07-OCT-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/952,545
FILING DATE: 30-SEP-1992
ATTOCNEY/AGENT INFORMATION:
NAME: Ihnen, Jeffrey L.
REGISTRATION NUMBER: 28,957
REFERNCE/DOCKET NUMBER: 23.33-124
TELEPHONE: 202-783-6040
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "primer"
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Patent No. 6165727
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFRUNE: 202-783-6031
INFORMATION FOR SEQ ID NO: 19: SEQUENCE CHARACTERISTICS: LENGTH: 20 base pairs TYPE: nucleic acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        18 ccgacciccaciccaci
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Washington
              D.C.
USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-09-106-216-19
                                                   COUNTRY:
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PRIOR APPLICATION NUMBER:
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PAVIRANI, ANDREA
PERRAUD, FREDERIC
VENTION: CELL LINEAGES EXPRESSING A BIOLOGICALLY
VENTION: ACTIVE IX FACTOR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-632-575B-21/c

Sequence 21, Application US/08632575B
Fatent No. 5843660
GENERAL INFORMATION:
APPLICANT: Schumm, James W.
TITLE OF INVENTION: Multiplex Amplification of
TITLE OF INVENTION: Multiplex Amplification of
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSEE: 2800 Woods Hollow Road
STREET: 2800 Woods Hollow Road
                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: PROMPH COMPATIBLE
COMPUTER: PROMPH COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/394,210
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 2800 mcc...
STREET: 2800 mcc...
CITY: Madison
STATE: Wisconsin
COUNTRY: U.S.A.
ZIP: 53711-539
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 16.2; DB 1;
Pred. No. 5.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: PEW/5683/84493
TELECOMMUNICATION INFORMATION:
TELEPHONE: 202-861-3000
TELEFAC: (202) 861-0944
TELEX: 6714627 CUSH
INFORMATION FOR SEQ ID NO: 6:
ERGURNCE CHARACTERISTICS:
LENGTH: 21 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                         CLASSIFICATION: 800
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US/08/038,085
FILING DATE: 09-APR-1991
APPLICATION NUMBER: FR 8910720
FILING DATE: 09-APR-1991
APPLICATION NUMBER: FR 8910720
FILING DATE: 09-AUG-1989
ATTORNEY/AGENT INFORMATION:
NAME: WHITE JR, PAUL E
REGISTRATION NUMBER: 32011
                                                                         CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2352 GATTACAGGCATGAGCCACCG 2372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                GATTATAGGCGTGAGCCACTG 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity 85.7%;
Matches 18; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                         NUMBER OF SEQUENCES: 2
CORRESPONDENCE ADDRESS:
ADDRESSEE: CUSHMAN,
                                     TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                           COUNTRY:
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Sequence 80, Application US/09157177

Patent No. 609058

GENERAL INPORMATION:
APPLICANT: Butler, John M.
APPLICANT: Li, Jia
APPLICANT: Butler, Joseph A.
APPLICANT: Butler, Joseph A.
APPLICANT: Becker, Christopher H.
TITLE OF INVENTION: REPEAT MARKERS
TITLE OF INVENTION: REPEAT MARKERS
TITLE OF INVENTION: REPEAT MARKERS
CURRENT APPLICATION NUMBER: US/09/157,177

CURRENT FILING DATE: 1998-09-18

EARLIER PRILING DATE: 1997-09-19

NUMBER OF SEQ ID NOS: 135

CONTWARE PATENTION OF: 2.0
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Sequence 21, Application US/09199542B

Patent No. 647921S

GENERAL INFORMATION

APPLICANT: Schumm, James W.

TITLE OF INVENTION: Multiplex Amplification of Short Tandem Repeat Loci
FILE REFERENCE: 16026/9212

CURRENT APPLICATION NUMBER: US/09/199,542B

CURRENT FILING DATE: 1998-11-25
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0.7%; Score 16.2; DB 1; Length 21;
Best Local Similarity 85.7%; Pred. No. 5.3e+02;
Matches 18; Conservative 0; Mismatches 3; Indels
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Pred. No. 5.3e+02;
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SYSTEM: DOS, version 6.0
WordPerfect 5.1 (DOS text format)
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Best Local Similarity 85.7%; Pred. No. 5.3e
Matches 18; Conservative 0; Mismatches
       CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/632,575B
FILING DATE: 04/15/96
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/316,544
FILING DATE: 09/30/94
; INFORMATION FOR SEQ ID NO: 21:
SEQUENCE CHARACTERISTICS:
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POSITION IN GENOWE:

MAP POSITION: D14S548

US-08-632-5758-21
                                                                                                                                                                                                                                                            LENGTH: 21
TYPE: Nucleic Acid
STRANDEDNESS: Single
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CORGANISM: Homo sapiens
US-09-157-177-80
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Butler, John M.
APPLICANT: Butler, John M.
APPLICANT: Butler, John M.
APPLICANT: Butler, John M.
APPLICANT: Butler, Joseph A.
APPLICANT: Monforte, Joseph A.
APPLICANT: Becker, Christopher H.
TITLE OF INVENTION: DNA TYPING BY MASS SPECTROMETRY WITH POLYMORPHIC DNA
TITLE OF INVENTION: DNA TYPING BY MASS
FILE REFERENCE: GETR: 0.17/GETR017P;
CURRENT APPLICATION NUMBER: US/09/541,210
CURRENT FILING DATE: 1997-09-19
ARALIER PILING DATE: 1997-09-19
NUMBER OF SEQ ID NOS: 135
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 80
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Sequence 91, Application US/09347114A

Patent No. 6297014

GENERAL INFORMATION:

APPLICANT: Maren T. Scheuner (Inventor)

APPLICANT: Maren T. Scheuner (Inventor)

APPLICANT: Huiying Yang (Inventor)

TITLE OF INVENTION: No. 6297014-responsiveness to Statin Drug Treatment

FILE REFERENCE: P07 41878

CURRENT APPLICATION: NO. 6297014-responsiveness to Statin Drug Treatment

FILE REFERENCE: P07 41878

CURRENT APPLICATION: NUMBER: US/09/347,114A

CURRENT FILING DATE: 1999-07-02

NUMBER OF SEQ ID NOS: 110

SEQ ID NO 91

LENGTH: 17
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                                                                                                                                                                                                                                                                                                                                  3; Indels
                                                                                                                                                                                                                                                                                  0.7%; Score 16.2; DB 1;
85.7%; Pred. No. 5.3e+02;
tive 0; Mismatches 3;
PRIOR FILING DATE: 1994-09-30
PRIOR APPLICATION NUMBER: US 08/632,575
PRIOR FILING DATE: 1996-04-15
NUMBER OF SEQ ID NOS: 110
SOFTWARE: Word97 (converted to DOS text format)
LENGTH: 21
                                                                                                                                                                                                                                                                                                                                                                           2107 AGTCTTGCTCTGTTACCCAGG 2127
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Best Local Similarity 85.73
Matches 18; Conservative
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; ORGANISM: Homo sapiens
US-09-541-210-80
                                                                                                                                                                 ; TYPE: DNA
; ORGANISM: Homo sapien
; LOCATION: D14S548
US-09-199-542B-21
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APPLICANT: Penny, Laura
APPLICANT: Galvin, Margaret
APPLICANT: Galvin, Margaret
APPLICANT: Galvin, Margaret
APPLICANT: Galvin, Margaret
APPLICANT: Michael
TITLE OF INVENTION: Genctyping Human
TITLE OF INVENTION: UPF-Glucuronosyltransferase 2B4 (UGT2B4), 2B7 (UGT2B7) and
TITLE OF INVENTION: 2B15 (UGT2B15) Genes
FILE REPERENCE: SEQ-22PRV2
CURRENT APPLICATION NUMBER: US/09/356,806
CURRENT FILING DATE: 1999-07-20
NUMBER OF SEQ ID NOS: 164
SOFTWARE: FastSEQ for Mindows Version 3.0
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APPLICANT: Lessard, Philip A.
APPLICANT: Lessard, Philip A.
APPLICANT: wills, Laura B.
APPLICANT: wills, Laura B.
APPLICANT: Stephanopoulos, Gregory
TITLE OF INVENTION: Pyruvate Carboxylase from Corynebacterium glutamicum
FILE REFERENCE: 1533.0790000
CURRENT APPLICATION NUMBER: US/09/220,081
CURRENT FILING DATE: 1998-12-23
NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 34
LENGTH: 19
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US-09-220-081-34
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                                                                        Length 17;
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                                                                                                                    0; Indels
                                                                      0.7%; Score 16; DB 1; Le
100.0%; Pred. No. 5.8e+02;
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100.0%; Pred. No.
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Patent No. 6586175
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 34, Application US/09220081 Patent No. 6171833 GENERAL INFORMATION:
                                                                                                                                                             2118 GTTACCCAGGCTGGAG 2133
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2274 GGGTTTCACCGTGTTA 2289
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                      16 GTTACCCAGGCTGGAG 1
                                                                                         Best Local Similarity 100.0
Matches 16; Conservative
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Best Local Similarity 100.(
Matches 16; Conservative
; ORGANISM: Homo sapiens
US-09-347-114A-91
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CORGANISM: H. sapiens
US-09-356-806-13
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LENGTH: 17
                                                                 Query Match
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Wed Jan 26 12:47:30 2005

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PRIOR FILING DATE: 1998-12-23
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LENGTH: 19
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Patent No. 6403351
GENERAL INCRMATION:
GENERAL INCRMATION:
APPLICANT: Sinakey, Anthony J.
APPLICANT: Willis, Laura B.
APPLICANT: Willis, Laura B.
TILLE OF INVENTION: Pyruvate Carboxylase from Corynebacterium glutamicum FILE REFERENCE: 1533.079000
CURRENT APPLICATION NUMBER: US/09/677,575
CURRENT FILING DATE: 2000-10-03
PRIOR APPLICATION NUMBER: 09/220,081
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              0.7%; Score 16; DB 1; Length 19; 100.0%; Pred. No. 5.6e+02;
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                                                                                 US-09-366-840-2

; Sequence 2, Application US/09366840
; Patent No. 6228145;
; GENERAL INCEMATION:
; APPLICANT: OSSOWSKi, Liliana
; TITLE OF INVENTION: In Vivo Assay for Intravasation
; FILE REFERENCE: A32590 70165.0550
; CURRENT APPLICATION WUMBER: US/09/366,840
; CURRENT FILING DATE: 1999-08-04
; SOFTWARE: PatentIn Ver. 2.1
; SEQ ID NO 2
; LENGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2, Application US/09078294
Patent No. 626511
GENERAL INFORMATION:
APPLICANT: Choo, Kong-Hong Andy
APPLICANT: Choo, Michael R.
TITLE OF INVENTION: A NOVEL NUCLEIC ACID MOLECULE
FILE REFERENCE: Davies Col
CURRENT PLILNG DATE: 1998-05-13
NUMBER OF SEQ ID NOS: 29
SOCTWARE: PatentIN Ver. 2.0
SEQ ID NO SEQ ID NOS: 29
LENGTH: 19
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  4 GTGGAATGATCCCCGA 19
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ORGANISM: DNA primer
US-09-078-294-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 16; Conserv
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; ORGANISM: Human
US-09-366-840-2
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Gaps
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Patent No. 6165786
GEBREAL INFORMATION:
APPLICANT: C. Frank Bennett
TITLE OF INVENTION: ANTISENSE MODULATION OF NUCLEOLIN EXPRESSION
FILE REPERENCE: RTS-0109
CURRENT APPLICATION NUMBER: US/09/433,699B
CURRENT FILING DATE: 1999-11-03
SEQ ID NO 43
LENGTH: 20
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                                                                                                                                                                                                                                                         Query Match 0.7%; Score 16; DB 1; Length 19; Best Local Similarity 100.0%; Pred. No. 5.6e+02; Matches 16; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 222. Application US/09544398B

Patent No. 6770461

GENERAL INFORMATION:
APPLICANT: Carulli, John P.
APPLICANT: Carulli, John P.
APPLICANT: Recker, Robert R.
APPLICANT: Recker, Robert R.
TITLE OF INVENTION: High bone mass gene of 11q13.3
FILE REFERENCE: 032796.013
CURRENT APPLICATION WUMBER: US/09/544,398B
CURRENT APPLICATION WUMBER: US 09/229,319
PRIOR FILING DATE: 1999-01-13
PRIOR APPLICATION NUMBER: US 60/071,449
PRIOR FILING DATE: 1999-01-13
PRIOR PLING DATE: 1999-01-23
PRIOR PLING DATE: 1999-10-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 641.
SOFTWARE: FastSEQ for Windows Version 4.0
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                                                                                                                                                                                                                                                                                                                                                                    185 GTGGAATGATCCCCGA 200
                                                                                                        TYPE: DNA
ORGANISM: Artificial Sequence
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NUMBER OF SEQ ID NOS: 36
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 34
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Sequence 11, Application US/08757223

Sequence 11, Application US/08757223

Patent No. 6136530

GENERAL INFORMATION:

APPLICANT: Poduslo, Shirley E.

TITLE OF INVENTION: COMPOSITIONS AND METHODS FOR ASSESSING RISK

TITLE OF INVENTION: FACTORS IN ALZHEIMER'S DISEASE

NUMBER OF SEQUENCES: 13

CORRESPONDENCE ADDRESS:

ADDRESSEE: Locke Purnell Rain Harrell

STREET: 2200 Ross Avenue, Suite 2200

CITY: Dallas

STATE: Texas

STATE: Texas

STATE: HBM PC compatible

COMPUTER: READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: BM PC compatible

COMPUTER: PATENTIN RC-DOS/MS-DOS

SOFTWARE: PATENTIN RC-DOS/MS-DOS

SOFTWARE: PATENTIN RC-DOS/MS-DOS

CURRENT APPLICATION NUMBER: US/08/757,223

PILING DATE: NO. 6136530ember 27, 1996
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APPLICANT: DIETWALER, Wolfgang
APPLICANT: FISHEL, Richard
TITLE OF INVENTION: METHOD FOR THE DETETION OF MICROSATELLITE
TITLE OF INVENTION: INSTABILLITY FOR TUMOR DIAGNOSTICS
NUMBER OF SEQUENCES: 50
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.7%; Score 15.8; DB 1; Length 19; Best Local Similarity 89.5%; Pred. No. 5.8e+02; Matches 17; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Nikaido, Marmelstein, Murray & Oram LLP STREET: 655 Fifteenth Street N.W. Suite 330 CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Patentin Release #1.0, Version #1.30
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APPLICATION NUMBER: US/09/047,347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2342 AAAGTGCTGGGATTACAGG 2360
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Patent No. 6150100
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Mayfield, Denise L.
REPERENCE/DOCKET NUMBER: 4-00
TELECOMMUNICATION INFORMATION:
TELEPHONE: 214/740-8785
TELEFRAX: 214/740-8800
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 11:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  19 AAAGTGGTGGGGTTACAGG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TOPOLOGY: linear
US-08-757-223-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3: D.C.
RY: U.S.A.
20005-5701
                              RESULT 612
US-08-757-223-11/c
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COUNTRY:
ZIP: 2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 613
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       8
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US-09-467-642-64/C

Sequence 64, Application US/09467642

Sequence 64, Application US/09467642

Sequence 64, Application US/09467642

GENERAL INFORMATION:

APPLICANT: Lex M. Cowment

APPLICANT: Lex M. Cowment

TILLE DE INVENTION:

FILLE REFERENCE: RTS-01.06

CURRENT APPLICATION NUMBER: US/09/467,642

CURRENT PILING DATE: 1999-12-20

NUMBER OF SEQ ID NOS: 89

SEQ ID NO 64

LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:

APPLICANT: Buller, John M.
APPLICANT: Monforte, Joseph A.
APPLICANT: Monforte, Joseph A.
TITLE OF INVENTION: BREBEAT WARKERS
FILE OF INVENTION: REPEAT WARKERS
FILE REFERENCE: GETR: 017/GETR017P
CURRENT APPLICATION NUMBER: 05/09/157,177
CURRENT FILLING DATE: 1998-09-18
EARLIER APPLICATION NUMBER: 60/059,415
BARLIER FILING DATE: 1997-09-19
NUMBER OF SEQ ID NOS: 135
SEQ ID NO 74
SEQ ID NO 74
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89.5%; Pred. No. 5.8e+02;
live 0; Mismatches 2; Indels
                                                                             Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 20;
                                                                                                                          Indels
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                                                                        Ouery Match
0.7%; Score 16; DB 1; L.
Best Local Similarity 100.0%; Pred. No. 5.5e+02;
Matches 16; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
0.7%; Score 16; DB 1; Lu
Best Local Similarity 100.0%; Pred. No. 5.5e+02;
Matches 16; Conservative 0; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FEATURE:

, OTHER INFORMATION: Antisense Oligonucleotide

US-09-467-642-64
OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-157-177-74/c
; Sequence 74, Application US/09157177
; Patent No. 6090558
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                                                                                                                                                                       1119 GATGAAGATGATGAGG 1134
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19 GAGTCTTGCTCTGTCGCCC
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Best Local Similarity
Matches 17; Conserv
          ; OTHER INFORM
US-09-433-699-43
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SOFTWAIRS PARENTE RELOGINGS OF THE SOFTWAIRS PARENT APPLICATION DATA:

APPLICATION NUMBER: US/09/402,923A
FILING DATE: 14-Feb-2001
PRIOR APPLICATION DATA:

RAPLICATION NUMBER: CT/(GB98/01102
FILING DATE: 15-ARR-1998
APPLICATION NUMBER: US 60/043,553
FILING DATE: 15-ARR-1997
APPLICATION NUMBER: US 60/048,740
FILING DATE: 05-JUN-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSER: Nixon and Vanderhye
STREET: 1100 No. 6555654th Glebe Road, Eighth Floor
CITY: Arlington
STATE: Virginia
COUNTRY: US
ZIP: VA 22201-4714
COMPUTER READABLE FORM:
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: IBM PC COMPATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                Length 19;
                                                                                                                                                                                                                                                                                                                                                                             2; Indels
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Merriman, Tony R
Merriker, Michael L
TITLE OF INVENTION: No. 6555654el LDL-Receptor
NUMBER OF SEQUENCES: 455
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                          Query Match 0.7%; Score 15.8; DB 1; Best Local Similarity 89.5%; Pred. No. 5.8e+02; Matches 17; Conservative 0; Mismatches 2;
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SEQUENCE DESCRIPTION: SEQ ID NO: 242:
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NAME: B.J. Sadoff
REGISTRATION NUMBER: 36,663
REFERENCE/DOCKET NUMBER: 620-81
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 615
US-09-402-923A-242/c
US-09-402-923A-242/c
Sequence 242, Application US/09402923A
Patent No. 6555654
GENERAL INFORMATION:
APPLICANT: Todd, John A
Hess, John W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                       2234 CACCACACCTGGCTAATTT 2252
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19 caccarcccrcccrcarrr 1
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Gerhold, David
Hammond, Holly
                                                                242:
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SEQUENCE CHARACTERISTICS:
TRNGTH: 19 hase pairs
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STRANDEDNESS: single
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                                                      INFORMATION FOR SEQ ID NO: 2 SEQUENCE CHARACTERISTICS: 1 LENGTH: 19 base pairs 1 TYPE: nucleic acid 1 STRANDEDNESS: single 1 TOPOLOGY: linear US-09-060-299-242
                                (703)816-4100
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                                TELEFAX:
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/060,299
FILING DATE: 15-APR-1998
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/043,553
FILING DATE: 15-APR-1997
PRIOR APPLICATION NUMBER: US 60/048,740
APPLICATION NUMBER: US 60/048,740
ATORNEY/AGENT INFORMATION:
NAME: 05-JUN-1997
ATORNEY/AGENT INFORMATION:
NAME: 05-JUN-1997
ATORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Hadmuch, Holly
APPLICANT: Hawguchi, Yoshihiko
APPLICANT: Kawaguchi, Yoshihiko
APPLICANT: Merziman, Tony R
APPLICANT: Merzker, Michael L
TITLE OF INVENTION: No. 6545137el Receptor
NUMBER OF SEQUENCES: 455
CORRESPONDENCE ADDRESS:
ADDRESSEE: 1100 No. 6545137th Glebe Road, Eighth Floor
CITY: Arlington
STATE: Virginia
COUNTRY: US
CONPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
CORTANDED: DAILORS COUNTRY: OF COMPUTER: OPERATING SYSTEM: PC-DOS/MS-DOS
CORTANDED: DAILORS COUNTRY: OF COMPUTER: OPERATING SYSTEM: PC-DOS/MS-DOS
CORTANDED: DAILORS COUNTRY: OF COMPUTER: OPERATING SYSTEM: PC-DOS/MS-DOS
COUNTRY: OPERATING SYSTEM: PC-DOS/MS-DO
REGISTRATION NUMBER: 36,663
REPERENCE/DOCKET NUMBER: 620-35
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)816-4091
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 242, Application US/09060299
Patent No. 6545137
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APPLICANT: Fodd, John A
APPLICANT: Hess, John W
APPLICANT: Caskey, Charles T
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cox, Roger D
Gerhold, David
Hammond, Holly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity 89.5°
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: DNA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
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Query Match
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                                                                                                                                                                                                                                                                      GENERAL INFORMATION:
APPLICANT: Butler, John M.
APPLICANT: Butler, John M.
APPLICANT: Li, Jia
APPLICANT: Li, Jia
APPLICANT: Bocker, Christopher H.
TITLE OF INVENTION: DNA TYPING BY MASS SPECTROMETRY WITH POLYMORPHIC DNA
TITLE OP INVENTION: REPEAT MARKERS
FILE REFERENCE: GETR: 0.17/GETR0.17p
CURRENT APPLICATION NUMBER: US/09/541,210
CURRENT APPLICATION NUMBER: 60/059,415
EARLIER APPLICATION WINGHER: 60/059,415
EARLIER FILING DATE: 1997-09-19
NUMBER: OF SEQ ID NOS: 135
SOFID NO 74
SEC ID NO 74
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APPLICANT: Robbins, Joan M.
APPLICANT: TITLE, Richard
TITLE OF INVENTION: SKIN AND BYE DISEASES
THE REPERRICE: 480124.407
CURRENT APPLICATION NUMBER: US/09/696,791
CURRENT FILING DATE: 2000-10-25
NUMBER OF SEQ ID NOS: 4523
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 3179
LENGTH: 19
                                                    Gaps
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      0.7%; Score 15.8; DB 1; Length 19; 89.5%; Pred. No. 5.8e+02; ive 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 15.8; DB 1; Length 19;
Pred. No. 5.8e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Cyclin Al ribozyme binding site US-09-696-791-3179
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 15.8; DB 1;
Pred. No. 5.8e+02;
0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 3179, Application US/09696791 Patent No. 6770633
                                                                                                                                                                                                                                          Sequence 74, Application US/09541210
Patent No. 6764822
                                                                                       2234 CACCACACCTGGCTAATTT 2252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2106 GAGICITGCTCTGTTACCC 2124
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                                                                                                                                 19 caccardccrddcraarrr 1
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0.7%;
Best Local Similarity 89.5%;
Matches 17; Conservative 0
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Best Local Similarity 89.5%;
Matches 17; Conservative
Query Match 0.74
Best Local Similarity 89.5'
Matches 17; Conservative
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CAGANISM: Homo sapiens
US-09-541-210-74
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA
ORGANISM: Homo sapiens
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US-09-696-791-3179/c
                                                                                                                                                                                                                      US-09-541-210-74/C
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APPLICANT: Robbins, Joan M.
APPLICANT: Robbins, Joan M.
APPLICANT: Tritz, Richard
TITLE OF INVENTION: RIBOZYME THERAPY FOR THE TREATMENT OF PROLIFERATIVE
TITLE OF INVENTION: SKIN AND EYE DISEASES
FILE REFERENCE: 480124.407
CURRENT PAPLICATION NUMBER: US/09/696,791
CURRENT FILING DATE: 2000-10-25
NUMBER OF SEQ ID NOS: 4523
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 3630
LENGTH: 19
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APPLICANT: Tritz, Richard
TITLE OF INVENTION: RIBOZIME THERAPY FOR THE TREATMENT OF PROLIFERATIVE
TITLE OF INVENTION: SIN AND EYE DISEASES
FILE REFERENCE: 480124.407
CURRENT APPLICATION NUMBER: US/09/696,791
CURRENT FILING DATE: 2000-10-25
NUMBER OF SEQ ID NOS: 4523
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 3631
LENGTH: 19
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Patent No. 6770633
GENERAL INFORMATION
APPLICANT: Robbins, Joan M.
APPLICANT: Tritz, Richard
TITLE OF INVENTION: RIBOZYME THERAPY FOR THE TREATMENT OF PROLIFERATIVE
TITLE OF INVENTION: SKIN AND EYE DISEASES
FILE REPERENCE: 480124.407
CURRENT APPLICATION NUMBER: US/09/696,791
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89.5%; Pred. No. 5.8e+02;
cive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; OTHER INFORMATION: Cdc25 hs ribozyme binding site US-09-696-791-3630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Cdc25 hs ribozyme binding site US-09-696-791-3631
                    Sequence 3630, Application US/09696791
Patent No. 6770633
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Patent No. 6770633
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Matches 17; Conservative
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ORGANISM: Homo sapiens
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ORGANISM: Homo sapiens
                                                                                  GENERAL INFORMATION:
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18-09-696-791-3630
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0.7%; Score 15.8; DB 1; Length 20;
Best Local Similarity 89.5%; Pred. No. 5.7e+02;
Matches 17; Conservative 0; Mismatches 2; Indels
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Mocdecock Washburn Kurtz
ADDRESSEE: Mocdecock Washburn Kurtz
ADDRESSEE: Mocdecock Washburn Kurtz
ADDRESSEE: Mackiewicz & No. 5656743ris
STREET: One Liberty Place - 46th Floor
CITY: Philadelphia
STATE: 0.0 Liberty Place - 46th Floor
CITY: Philadelphia
STATE: 0.0 Liberty Place - 46th Floor
CUNTRY: USA
ZAPE: Diskette, 3.5 inch, 1.44 Mb stor.
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.5 inch, 1.44 Mb stor.
COMPUTER: Diskette, 3.5 inch, 1.44 Mb stor
                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Cdc25 hs ribozyme binding site US-09-696-791-3632
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Patent No. 5656743
CENERAL INFORMATION:
APPLICANT: Busch et al.
TITLE OF INVENTION: OLIGONUCLEOTIDE MODULATION
TITLE OF INVENTION: OF CELL GROWTH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   817 AAGAAATTCAGATGAATT 835
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CURRENT FILING DATE: 2000-10-25
NUMBER OF SEQ ID NOS: 4523
SOFTWARE: PatentIn Ver. 2.0
SEQ ID NO 3632
LENGTH: 19
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STRANDEDNESS: single
                                                                                                                                                                                                             TYPE: DNA
ORGANISM: Homo sapiens
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2232 GCCACCACCTGGCTAAT 2250

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APPLICANT: Schum, James W.
APPLICANT: Schum, James W.
APPLICANT: Bacher, Jeffery W.
TITLE OF INVENTION: MATERIALS AND METHODS FOR
TITLE OF INVENTION: IDENTIFYING AND ANALYZING INTERMEDIATE TANDEM
TITLE OF INVENTION: REPEAT DNA MARKERS
NUMBER OF SEQUENCES: 147
CORRESPONDENCE ADDRESS:
ADDRESSEE: Promega Corporation
STREET: 2800 Woods Hollow Road
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Sequence 164, Application US/09289267A

Sequence No. 6046320
GENERAL INFORMATION:
GENERAL INFORMATION:
THOUSE OF INVENTION: ANTISENSE MODULATION OF MDMX EXPRESSION
TITLE OF INVENTION: ANTISENSE MODULATION OF MDMX EXPRESSION
TITLE REFERENCE: RTS-0049
CURRENT APPLICATION NUMBER: US/09/289,267A
NUMBER OF SEQ ID NOS: 166
SEQ ID NO 164
LENGTH: 20
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0.7%; Score 15.8; DB 1;
Best Local Similarity 89.5%; Pred. No. 5.7e+02;
Matches 17; Conservative 0; Mismatches 2;
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CITY: Madison
STATE: Wisconsin
COUNTY: U.S.A.
ZIP: 5311-5399
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb
COMPUTER: IBM compatible PC
OPERATING SYSTEM: Windows 95
SOFTWARE: Word 97 (DOS text format)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/018,584A
FILING DATE: 04-Feb-1998
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Grady J. Frenchick
REGISTRATION NUMBER: 29,018
REGISTRATION NUMBER: 29,018
REGISTRATION NUMBER: 16026.9180
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Antisense Oligonucleotide US-09-289-267-164
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; Sequence 120, Application US/09018584A
; Patent No. 6218863
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   2127 GCTGGAGTGCAGTGGGTGA 2145
                                                                                                                                                                                                                                                                                                                                                        TYPE: DNA ORGANISM: Artificial Sequence
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STRANDEDNESS: Single
TOPOLOGY: Linear
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APPLICANT: Breet P. Monia

APPLICANT: William Gaarde

APPLICANT: William Gaarde

APPLICANT: Susan M. Freier

APPLICANT: Edward V. Wancewicz

TITLE OF INVENTION: ANTISENSE MODULATION OF TERT EXPRESSION

FILE REFERENCE: ISPH-0527,

CURRENT APPLICATION NUMBER: US/09/733,294A

CURRENT FILING DATE: 2000-12-07

PRIOR APPLICATION NUMBER: 09/572,423

PRIOR FILING DATE: 2000-05-16

NUMBER OF SEQ ID NOS: 108
APPLICANT: Hong Zhang
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF APAF-1 EXPRESSION
FILE REPRENCE: RTS-0190
CURRENT APPLICATION NUMBER: US/09/690,364
CURRENT FILING DATE: 2000-10-17
NUMBER OF SEQ ID NOS: 100
LENGTH: 20
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89.5%; Pred. No. 5.7e+02;
tive 0; Mismatches 2;
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0.7%; Score 15.8; DB 1;
Best Local Similarity 89.5%; Pred. No. 5.7e+02;
Matches 17; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 83, Application US/09918686; Patent No. 6475739; GENERAL INFORMATION: APPLICANT: Brunkow, Mary; APPLICANT: Brunkow, Mary; APPLICANT: Brench Sean, APPLICANT: Reaper, Bryan; APPLICANT: Stachling-Hampton, Karen; TITLE OF INVENTION: GENOMIC DELETIONS; FILE REPRENCE: 240083.515; CURRENT APPLICATION NUMBER: US/09/918,686; CURRENT FILING DATE: 2001-07-30; NUMBER OF SEQ ID NOS: 105; SOFTWARE: FastSEQ for Windows Version 4.0; SEQ ID NO 83.
                                                                                                                                                                                                                                                                                                                                                                      ; FEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-09-690-364-48
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                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Artificial Sequence
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Best Local Similarity 89.5'
Matches 17; Conservative
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US-09-733-294A-75
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Retent No. 6300132

GENERAL INFORMATION:
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GENERAL INFORMATION:
APPLICANT:
LEX M. COWBERT
TILLE OF INVENTION:
APPLICATION WOMBER:
US/09/467,642
UNRENT FILING DATE:
UNMER OF SEQ ID NOS:
89
LENGTH: 20
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Batent No. 6410334
GENERAL INFORMATION:
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APPLICANT:
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                            0.7%; Score 15.8; DB 1; Length 20; 89.5%; Pred. No. 5.7e+02; tive 0; Mismatches 2; Indels
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89.5%; Pred. No. 5.7e+02;
tive 0; Mismatches 2; Indel8
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                                                                                                                                                                 2099 TGAGACCGAGICTIGCICT 2117
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; Sequence 48, Application US/09690364
; Patent No. 6468795
; GENERAL INFORMATION:
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                        Query Match
Best Local Similarity 89.5'
Matches 17; Conservative
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Best Local Similarity 89.5
Matches 17; Conservative
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US-09-467-642-63/c
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EPO)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Nixon and Vanderhye
STREET: 1100 No. 6555654th Glebe Road, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2; Indels
                     PatentIn Release #1.0, Version #1.25 (EPO)
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TITLE OF INVENTION: No. 6555654el LDL-Receptor
NUMBER OF SEQUENCES: 455
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.7%; Score 15.8; DB 1;
89.5%; Pred. No. 5.7e+02;
tive 0; Mismatches 2;
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APPLICATION UNDER: US/09/402,923A
FILING DATE: 14-Feb-2001

PRIOR APPLICATION DATA:
APPLICATION NUMBER: PCT/GB98/01102
FILING DATE: 15-APR-1998
APPLICATION NUMBER: US 60/043,553
FILING DATE: 15-APR-1997
                                                        APPLICATION NUMBER: US/09/060,299
FILING DATE: 15-APR-1998
CLASSIPICATION 435
FRIOR APPLICATION DATA:
APPLICATION NUMBER: US 60/043,553
FILING DATE: 15-APR-1997
PRIOR APPLICATION NUMBER: US 60/048,740
FILING DATE: 05-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAMME: B.J.Sadoff
REGISTRATION NUMBER: 36,663
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 296, Application US/09402923A
Patent No. 655654
GENERAL INFORMATION:
APPLICANT: Todd, John A
Hess, John W
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Kawaguchi, Yoshihiko
Merriman, Tony R
PC-DOS/MS-DOS
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Cox, Roger D
Gerhold, David
Hammond, Holly
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COUNTRY: US
ZIP: VA 22201-4714
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2151 GCTCACTGCAAGCTCTGCC 2169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GTTCACTGCAACCTCTGCC 19
                                                                                                                                                                                                                                                                                                                                                                             TELEFAX: (703)816-4100
INFORMATION FOR SEQ ID NO: 296:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 62
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)816-4091
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hey, Patricia
                        SOFTWARE: PatentIn ReleCURRENT APPLICATION DATA: APPLICATION NUMBER: US,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 89.5
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Arlington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    nucleic acid
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APPLICANT: Hong Zhang
APPLICANT: Hong Zhang
APPLICANT: O'acqueline Wyatt
TITLE OF INVENTION: ANTIBENSE MODULATION OF BH3 INTERACTING DOMAIN DEATH AGONIST
TITLE OF INVENTION: EXPRESSION
FILE REFERENCE: RTS-0135
CURRENT APPLICATION NUMBER: US/09/657,346A
CURRENT FILING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 174
SEQ ID NO 52
LENGTH: 20
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                                                                                                                                                                    Length 20;
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1100 No. 6545137th Glebe Road, Eighth Floor
                                                                                                                                                               Query Match

0.7%; Score 15.8; DB 1;
Best Local Similarity 89.5%; Pred. No. 5.7e+02;
Matches 17; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match

0.7%; Score 15.8; DB 1;
Best Local Similarity 89.5%; Pred. No. 5.7e+02;
Matches 17; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ...cldANT: Hey, Patricia
APPLICANT: Kawaguchi, Yoshihiko
APPLICANT: Merriman, Tony R
APPLICANT: Merriman, Tony R
APPLICANT: Merzker, Michael L
TITLE OF INVENTION: NO. 6545137el Receptor NUMBER OF SEQUENCES: 455
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon
STREFT
                                                                          ; FEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-09-733-294A-75
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Antisense Oligonucleotide US-09-657-346A-52
                                                                                                                                                                                                                                                                                                                                                                                              Sequence 52, Application US/09657346A Patent No. 6503754 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 296, Application US/09060299
Patent No. 6543137
GENERAL INFORMATION:
APPLICANT: Todd, John A
                                                                                                                                                                                                                                                  2270 GACAGGGTTTCACCGTGTT 2288
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2254 TTGTACTTTAGTAGAGAC 2272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Arington
STATE: Virginia
COUNTRY: US
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                            1 GATAGGGTTTCACCATGTT 19
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Hess, John W
Caskey, Charles T
Cox, Roger D
Gerhold, David
Hammond, Holly
                                         TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TYPE: DNA ORGANISM: Artificial Sequence
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US-09-060-299-296
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APPLICANT:
SEQ ID NO 75
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RESULT 635
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; Sequence 73, Application US/09679299A
; Patent No. 6566135
; GENERAL INFORMATION:
    APPLICANT: Vickie L. Brown-Driver
; APPLICANT: Hong Zhang
    APPLICANT: Andrew T. Watt
    TITLE OF INVENTION: ANTISENSE MODULATION OF CASPASE 6 EXPRESSION
    TITLE OF INVENTION: ANTISENSE MODULATION OF CASPASE 6
    CURRENT APPLICATION NUMBER: US/09/679,299A
    CURRENT FILING DATE: 2000-10-04
    NUMBER OF SEQ ID NOS: 164
    SEQ ID NO 73
    LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-679-299A-69

Sequence 69, Application US/09679299A

Sequence 69, Application US/09679299A

Sequence 69, Application US/09679299A

GENERAL INFORMATION:

APPLICANT: Vickie L. Brown-Driver

APPLICANT: Hong Zhang

APPLICANT: Andrew T. Watt

TITLE OF INVENTION: AND SERVESSION

FILE REFERENCE: RTS-0187
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.7%; Score 15.8; DB 1; Length 20; Best Local Similarity 89.5%; Pred. No. 5.7e+02; Matches 17; Conservative 0; Mismatches 2; Indels
APPLICATION NUMBER: US 60/048,740
FILING DATE: 05-UN-1997
ATTORNEY/AGENT INFORMATION:
NAME: B.J.Sadoff
REGISTRATION NUMBER: 36,663
REFERENCE/BOCKET NUMBER: 620-81
TELECOMMUNICATION INFORMATION:
TELEPHONE: (703)816-4091
TELEPHONE: (703)816-4091
TELEPHONE: (703)816-4100
INFORMATION FOR SEQ ID NO: 296:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: NUCleic acid
STYPE: NUCLEic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Antisense Oligonucleotide US-09-679-299A-69
                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 296:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/09/679,299A CURRENT FILING DATE: 2000-10-04 NUMBER OF SEQ ID NOS: 164 SEQ ID NO 69 LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2151 GCTCACTGCAAGCTCTGCC 2169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2348 CTGGGATTACAGGCATGAG 2366
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                       US-09-402-923A-296
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REPEAT DNA MARKERS
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                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STATE: Wisconsin
COUNTRY: U.S.A.
ZIP: 53711-539
COMPUTER: READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb
COMPUTER: IBM compatible PC
OPETWARE: Word 97 (DOS text format)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/784,423
                                                                                                                                                                                                                                                                                                                                                                                                                      | GENERAL INCO. 6 / 10163
| GENERAL INFORMATION:
| APPLICANT: Buchardt, Ole
| APPLICANT: Buchardt, Ole
| APPLICANT: Buchardt, Michael
| APPLICANT: Nelsen, Peter E.
| APPLICANT: Nelsen, Peter E.
| APPLICANT: Berg, Rolf H.
| TITLE OF INVENTION: PEPTIDE NUCLEIC ACIDS SYNTHONS
| FILE PEFERENCE: ISPS-1999
| CURRENT APPLICATION NUMBER: US/08/468,719A
| CURRENT PILING DATE: 1995-06-06
| PRIOR APPLICATION NUMBER: US 08/108,591
| PRIOR FILING DATE: 1993-11-22
| NUMBER OF SEQ ID NOS: 48
| SEQ ID NOS: 48
| SEQ ID NO S: 40
| SEQ ID NO S: 40
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Bacher, Jeffery W.
TITLE OF INVENTION: MATERIALS AND METHODS FOR
                                                                                                                  Score 15.8; DB 1;
Pred. No. 5.7e+02;
0; Mismatches 2;
                                           ; OTHER INFORMATION: Antisense Oligonucleotide US-09-679-299A-73
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: Oligonucleotide Primer US-08-468-719A-40
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ADDRESSER: Promega Corporation
STREET: 2800 Woods Hollow Road
CITY: Madison
                                                                                                                                                                                                                                                                                                                                                                 US-08-468-719A-40; Sequence 40, Application US/08468719A; Patent No. 6710163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; Sequence 120, Application US/09784423
; Patent No. 6767703
; GENERAL INFORMATION:
                                                                                                                                                                                                                   2297 TGGTCTCGATCTCCTGACC 2315
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CTTA 2058
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ORGANISM: Artificial Sequence
                                                                                                                  Query Match
Best Local Similarity 89.5%;
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES:
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Sequence 61, Application US/08649950

Patent No. 640303

GENERAL INFORMATION:
APPLICANT: Shipman, Robert
APPLICANT: Leushner, James
APPLICANT: Leushner, James
TITLE OF INVENTION: METHOD AND REAGENTS FOR TESTING FOR
TITLE OF INVENTION: MITATIONS IN THE BRCA1 GENE
NUMBER OF SEQUENCES: 77
CORRESPONDENCE ADDRESS:
ADDRESSEE: Oppedahl & Larson
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              - 3.5 inch, 1.44 Mb Btorage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         B: Oppedahl & Larson
1992 Commerce Street Suite 309
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            OPERATING SYSTEM: MS DOS
SOFTWARE: Word Perfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/649,950
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2187 ATTCTCCTGCCTCAGCCTC 2205
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ZIP: 10598
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.
COMPUTER: IBM compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                 Query Match
Best Local Similarity 89.5
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              17; Conservative
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CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Best Local Similarity
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CITY: Yorktown
STATE: NY
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US-08-649-950-61/c
                             JS-08-332-766A-68
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 21
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: AUGUSTSTEE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: The PC compatible
COMPUTER: The PC compatible
COMPUTER: The PC compatible
COMPUTER: The PC COMPUTER: The PC COMPUTER: The PC COMPUTER: DOS WESTER: D
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| Sequence 68, Application US/08332766A
| Patent No. 2843847|
| Pat
FILING DATE: 15-Feb-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/018,584
FILING DATE: 04-Feb-1998
ATORNEY/AGENT INFORMATION:
NAME: Crady J. Frenchick
REGISTRATION NUMBER: 29,018
REFERENCE/DOCKET NUMBER: 16026.9180
TELEPHONE: (608) 257-3501
TELEPHONE: (608) 257-2275
INFORMATION FOR SEQ ID NO: 120
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
SEQUENCE DESCRIPTION: SEQ ID NO: 120
US-09-784-423-120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2099 TGAGACCGAGTCTTGCTCT 2117
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MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (202) 822-0944
TELEX: 6714627 CUSH
INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          21 base pairs
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STRANDEDNESS: single
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0.7%; Score 15.8; DB 1; Length 21;
89.5%; Pred. No. 5.6e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0.7%; Score 15.8; DB 1; Length 21;
89.5%; Pred. No. 5.6e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                Sequence 7, Application US/09357740

Faguence 7, Application US/09357740

Fatent No. 6348596

GENERAL INFORMATION:

APPLICANT: Lee, Linda G.

APPLICANT: Mullah, Khairuzzaman B.

APPLICANT: Mullah, Khairuzzaman B.

TITLE OF INVENTION: Francis T.

TITLE OF INVENTION: ASYMMETRIC CYANINE DYE QUENCHERS

FILE REFERENCE: 9584-007

CURRENT FILING DATE: 1999-07-20

EARLIER APPLICATION NUMBER: 09/012,525

KUMBER OF SEQ ID NOS: 22

SOFTWARE: Patentin Ver. 2.0

SEQ ID NOS: 22
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; LOCATION: 1..21
. OTHER INFORMATION: downstream amplification primer 99-3944 for SEQ 3423, in compleme
US-09-422-978-11288
                                                                                                                 APPLICANT: Cohen, Daniel
APPLICANT: Blumenfeld, Marta
APPLICANT: Blumenfeld, Marta
APPLICANT: Blumenfeld, Marta
APPLICANT: Blumesfor'. IJya
TITLE OF INVENTION: Biallelic markers for use in constructing a high density...
FILE REPERENCE: GENSET.020CP1
CURRENT APPLICATION NUMBER: US/09/422,978
CURRENT APPLICATION NUMBER: US 09/298,850
EARLIER APPLICATION NUMBER: US 60/109,732
FARLIER PILING DATE: 1999-04-21
FARLIER FILING DATE: 1998-11-23
EARLIER FILING DATE: 1998-11-23
FARLIER FILING DATE: 1998-04-21
NUMBER OF SEQ ID NOS: 11796
SEQ ID NO 11288
LENGTH: 21
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US-09-358-972-252
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.7%; Score 15.8; DB 1; Length 21; Best Local Similarity 89.5%; Pred. No. 5.6e+02; Matches 17; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Olson, Ryan J.
APPLICANT: Wood, Keith W.
APPLICANT: Welch, Roy
TITLE OF INVENTION: Nucleic Acid Detection
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILE REFERENCE: Pro-103 6669/75528
CURRENT APPLICATION NUMBER: US/09/358,972
CURRENT PILING DATE: 1999-07-22
EARLIER APPLICATION NUMBER: 09/252,436
EARLIER APPLICATION NUMBER: 09/252,436
EARLIER APPLICATION NUMBER: 09/042,287
EARLIER APPLICATION NUMBER: 09/042,287
                                                Sequence 11288, Application US/09422978
Patent No. 6537751
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 252, Application US/09358972
Patent No. 6235480
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1037 AGATCAGTTTAGTGTAGAA 1055
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GENERAL INFORMATION:
APPLICANT: Shultz, John W
APPLICANT: Lieppe, Donna
APPLICANT: Lieppe, Donna
APPLICANT: Mandrekar, Michelle
APPLICANT: Rophart, Daniel
APPLICANT: Rhodes, Richard B.
APPLICANT: Andrews, Christine A.
APPLICANT: Hartnett, James R.
APPLICANT: Hartnett, James R.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                19 AGATCAGTTGAGGGTAGAA 1
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ORGANISM: Artificial Sequence
FEATURE:
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SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 252
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Homo Sapiens
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: primer_bind
LOCATION: 1..21
                              1-09-422-978-11288/c
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APPLICANT: Cohen, Daniel

APPLICANT: Cohen, Daniel

APPLICANT: Cohen, Daniel

APPLICANT: Blumenfeld, Marta

APPLICANT: Chumakov, Ilya

TITLE OF INVENTION: Biallelic markers for use in constructing a high density...

FILE REPERENCE: GENSET.020CP1

CURRENT APPLICATION NUMBER: US/09/422,978

CURRENT PILING DATE: 1999-10-20

EARLIER FILING DATE: 1999-04-21

EARLIER PLILING DATE: 1999-04-21

EARLIER PILING DATE: 1998-04-21

MUMBER OF SEQ ID NOS: 11796

SEQ ID NO 10751
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           , OTHER INFORMATION: amplification primer for BRCA1 gene
US-08-649-950-61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.7%; Score 15.8; DB 1; Best Local Similarity 89.5%; Pred. No. 5.6e+02; Matches 17; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         RESULT 639
8-09-422-978-10751/c
; Sequence 10751, Application US/09422978
; Patent No. 6537751
                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: double
TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
HYPOTHETICAL: no
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1701 TGTGCAAAGAAGCTAAAGA 1719
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                                                                      NAME: Larson, Marina T.
REGISTRATION NUMBER: 32,038
REFERENCE/DOCKET NUMBER: VGEN
TELECOMMUNICATION INFORMATION:
TELEPHONE: (914) 245-3252
TELEFAX: (914) 962-4330
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    20 rérectaagaageraaaga 2
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                                                                                                                                                                                                                                                61:
                      FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Larson, Marina T.
                                                                                                                                                                                                                                                                                                                                                                                                                                    no
'PB: internal
                                                                                                                                                                                                                                        INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Homo Sapiens
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                      ANTI-SENSE: no
FRAGMENT TYPE: inte
ORIGINAL SOURCE:
ORGANISM: human
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ORGANISM: Homo (US-09-866-108A-7368)
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Patent No. 6686188

GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: APPLICANT: APANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: GHEN, Wensheng
APPLICANT: HANDON, Wark
ITITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
                                                 Gaps
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US-09-383-316-88
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    Length 17;
                                               1; Indels
Query Match

0.6%; Score 15.4; DB 1;
Best Local Similarity 94.1%; Pred. No. 6.3e+02;
Matches 16; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Wood, Keith W.
APPLICANT: Welch, Roy
TITLE OF INVENTION: Nacleic Acid Detection
FILE REFERENCE: PRO-104 6668/75529
CURRENT APPLICATION NUMBER: US/09/383,316
CURRENT FILING DATE: 1999-08-25
PRIOR APPLICATION NUMBER: 09/252,436
PRIOR FILING DATE: 1999-02-18
PRIOR FILING DATE: 1998-03-13
PRIOR FILING DATE: 1998-03-13
PRIOR FILING DATE: 1998-03-13
PRIOR FILING DATE: 1998-03-13
NUMBER OF SEQ ID NOS: 123
SOFTWARE: PATENTIN VET. 2.1
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                                                                                                                                                                                                                         US-09-383-316-88/c
; Sequence 88, Application US/09383316
: Patent No. 6391551
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2112 TGCTCTGTTACCCAGGC 2128
                                                                                       2112 TGCTCTGTTACCCAGGC 2128
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Andrews, Christine A. Hartnett, James R.
                                                                                                                                                                                                                                                                                                                                                                 Lieppe, Donna
Mandrekar, Michelle
Kephart, Daniel
Rhodes, Richard B.
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                                                                                                                                    17 recrerereacceage 1
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                                                                                                                                                                                                                                                                                                                     Shultz, John W
Lewis, Martin K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Olson, Ryan J.
Wood, Keith W.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 88
LENGTH: 17
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PRIOR FILING DATE: 2000-05-26

PRIOR PILING DATE: 2000-10-04

PRIOR PILING DATE: 2000-10-04

PRIOR PILING DATE: 2000-10-04

PRIOR PILING DATE: 2000-10-03-07

PRIOR PILING DATE: 2000-09-27

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30

PRIOR PELING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00669

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30
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OTHER INFORMATION: Description of Artificial Sequence:probe to Alu2
OTHER INFORMATION: human gene
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CURRENT APPLICATION NUMBER: US/09/790,417
CURRENT FILING DATE: 2001-02-22
PRIOR APPLICATION NUMBER: 09/358,972
PRIOR PLICATION NUMBER: 09/358,972
PRIOR PLICATION NUMBER: 09/358,972
PRIOR FILING DATE: 1999-01-21
PRIOR FILING DATE: 1999-03-13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Aeomica Sequence Listing Engine
Patent No. 6686188
SEQ ID NO 7368
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Patent No. 6730479
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Kephart, Daniel
Rhodes, Richard B.
Andrews, Christine A.
Hartnett, James R.
Gu, Trent
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ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Shultz, John W
APPLICANT: Lewis, Martin K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQ ID NOS: 290
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 252
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Best Local Similarity 94.1
Matches 16; Conservative
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Wood, Keith W.
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US-09-696-791-1869/c
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Patent No. 6770633
Face Trick INCOMATION:
APPLICANT: Tritz, Richard
TITLE OF INVENTION: SEIN AND EYE DISEASES
FILE REFERENCE: 480124.407
CURRENT FILING DATE: 2000-10-25
NUMBER OF SEQ ID NOS: 4523
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 1321
LENGTH: 19
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                                          Query Match 0.6%; Score 15.4; DB 1; Length 17; Best Local Similarity 94.1%; Pred. No. 6.3e+02; Matches 16; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 0.6%; Score 15.4; DB 1; Length 18; Best Local Similarity 94.1%; Pred. No. 6.2e+02; Matches 16; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match 0.6%; Score 15.4; DB 1; Length 19; Best Local Similarity 94.1%; Pred. No. 6.1e+02; Matches 16; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                         Sequence 438. Application US/09544398B
; Batent No. 6770461
; GENERAL INFORMATION:
; APPLICANT: Carulli, John P.
; APPLICANT: Carulli, John P.
; APPLICANT: Carulli, John P.
; APPLICANT: Little, Randall D.
; APPLICANT: Recker, Robert R.
; APPLICANT: Recker, Robert R.
; TITLE OF INVENTION: High bone mass gene of 11q13.3
; FILE REFREENCE: 032796-013
; CURRENT APPLICATION NUMBER: US/09/544,398B
; CURRENT FILING DATE: 2002-06-10
; PRIOR APPLICATION NUMBER: US 60/071,449
; PRIOR PILING DATE: 1998-01-13
; PRIOR FILING DATE: 1998-10-23
; NUMBER OF SEQ ID NOS: 641
; SOFTWARE: FastSEQ for Windows Version 4.0
; SEQ ID NO 4.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; OTHER INFORMATION: Cdk-we-hu ribozyme binding site
US-09-696-791-1321
                                                                                                                           2112 TGCTCTGTTACCCAGGC 2128
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      2345 GIGCIGGGATTACAGGC 2361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 Graciggarracagec 17
                                                                                                                                                         17 TGCTCTGTCACCCAGGC 1
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          , ORGANISM: Homo sapiens
US-09-544-398B-438
                                                                                                                                                                                                                                              US-09-544-398B-438
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 646
US-09-696-791-1321
US-09-790-417-252
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Sequence 2143, Application US/09696791

; Sequence 2143, Application US/09696791
; Patent No. 6770633
; GENERAL INFORMATION:
; APPLICANT: Robbins, Joan M.
; APPLICANT: Robbins, Tritz, Richard
; TITLE OF INVENTION: RIBOZYME THERAPY FOR THE TREATMENT OF PROLIFERATIVE
; TITLE OF INVENTION: SKIN AND EYE DISEASES
; TITLE OF INVENTION: US/09/696,791
; CURRENT APPLICATION NUMBER: US/09/696,791
; CURRENT FILING DATE: 2000-10-25
; NUMBER OF SEQ ID NOS: 4523
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 2143

LENGTH: 19
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                                                                                                                                                                Sequence 1869, Application US/09696791
Patent No. 6770633
GENERAL INFORMATION:
APPLICANT: Robbins, Joan M.
APPLICANT: Tritz, Richard
TITLE OF INVENTION: RIBOZYME THERAPY FOR THE TREATMENT OF PROLIFERATIVE
TITLE OF INVENTION: SKIN AND BYE DISEASES
FILE REFERENCE: 480124.407
CURRENT PELING DATE: 2000-10-25
NUMBER OF SEQ ID NOS: 4523
SOFTWARE PATENTIN Ver. 2.0
LENGTH: 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             , OTHER INFORMATION: Cyclin D1 ribozyme binding site US-09-696-791-1869
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match

0.6%; Score 15.4; DB 1;
Best Local Similarity 94.1%; Pred. No. 6.1e+02;
Matches 16; Conservative 0; Mismatches 1;
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US-08-564-002-25
; Sequence 25, Application US/08564002
; Patent No. 5714329
; GENERAL INFORMATION:
; APPLICANT: Dracopoli, Nicolas
1363 CACAAGCTGAAGAGGGC 1379
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1216 ATTGGAAATGCACTTCA 1232
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                                       CACAAGTTGAAGAGGC 19
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Homo sapiens
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SEQ ID NO 87
LENGTH: 20
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APPLICANT: Tucker, Margaret
APPLICANT: Goldstein, Alissa
TITLE OF INVENTION: Methods for the Diagnosis of a Genetic
TITLE OF INVENTION: Methods for the Diagnosis of a Genetic
TITLE OF INVENTION: Predisposition to Cancer Associated with Variant CDK4
TITLE OF INVENTION: Allele
NUMBER OF SEGUENCES:
ADDRESSEE: PLEHR, HOHBACH, TEST, ALBRITTON & HERBERT
STREET: A Embarcadero Center, Suite 3400
CITY: San Francisco
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ö
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Patent No. 6458532

GENERAL INFORMATION:

APPLICANT: Becra-Wadleigh, Sevilla D.
Gershon, Elliot S.
Badner, Judith A.
Goldin, Lynn R.
Berrettini, Wade H.
Yoshikawa, Takeo
Sanders, Alan R.
Esterling, Lisa Markers and Diagnostic
TITLE OF INVERTION: Chromosomal Markers and Diagnostic
TITLE OF INVERTION: Chromosomal Markers and Diagnostic
Tests for Manic-Depressive Illness
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.6%; Score 15.4; DB 1; Length 20; 94.1%; Pred. No. 5.9e+02; ve 0; Mismatches 1; Indels
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ADDRESSEE: Twonsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco
                                                                                                                                                                                                                                                            COUNTRY: USA
ZIP: 94111-4187
COMPUTER READABLE PORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-CONS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/564,002
                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION:
PILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: Sherwood, Pamela J.
REGISTRATION NUMBER: 36,677
REFERENCE/DOCKET NUMBER: A-62562
TELECOMMUNICATION INFORMATION:
TELEPAX: (415) 794-1989
TELEFXX: (415) 398-3249
INFORMATION FOR SEQ ID NO: 25:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 Dase pairs
"""" nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "Primer"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2339 CCCAAAGTGCTGGGATT 2355
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ZIP: 94111-3834
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 CCCAAAGTGCTGGAATT 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.6%;
Best Local Similarity 94.1%;
Matches 16; Conservative
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US-09-091-952A-152/c
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Gaps
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CURRENT APPLICATION UNMER: US/09/920,759
CURRENT FILING DATE: 2001-08-01
NUMBER OF SEQ ID NOS: 91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Indels
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                                                                                                                                                 PRIOR APPLICATION DATA:

APPLICATION NUMBER: US 60/029,278
FILING DATE: 28-0CT-1996
APPLICATION NUMBER: PCT/US97/19381
FILING DATE: 28-0CT-1997
ATTORNEY/AGENT INFORMATION:
NAME: Smith, Timothy L.
REGISTRATION NUMBER: 35,367
TELEGOMMUTCATION INFORMATION:
TELECOMMUTCATION INFORMATION:
TELECOMMUTCATION INFORMATION:
TELEFRAX: (415) 576-0300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ) LOCATION: 1...20
; COTHER INFORMATION: Clone 23 forward primer SEQUENCE DESCRIPTION: SEQ ID NO: 152: US-09-091-952A-152
                  OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/091,952A
FILING DATE: 19-Apr-1999
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0.6%; Score 15.4; DB 1;
94.1%; Pred. No. 5.9e+02;
tive 0; Mismatches 1;
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94.1%; Pred. No. 5.9e+02;
tive 0; Mismatches 1;
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COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 87, Application US/09920759; Patent No. 6537811; GENERAL INFORMATION:
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LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                            TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 152:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         3 Arcirectracia 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Artificial Sequence
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Best Local Similarity 94.11
Matches 16; Conservative
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APPLICANT: Susan M. Freier
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.6
Best Local Similarity 94.1
Matches, 16; Conservative
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Patent NO. 5273883
GENERAL INFORMATION
APPLICANT: Saiki, Randall K.
APPLICANT: Nasarabadi, Shanavaz L.
TITLE OF INVENTION: Methods and Reagents for G Gamma Globin TITLE OF INVENTION: Typing
NUMBER OF SEQUENCES: 58
CORRESPONDENCE ADDRESS: ROCHEMAIN-LA ROCHE INC.
STREET: 340 Kingsland Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.6%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 6.1e+02; Matches 17; Conservative 0; Mismatches 3; Indels
COMPUTER READABLE FORM:

MEDIUM TYPE: Diskette, 3.50 inch, 800 KD storage COMPUTER: Apple Macintosh oPERATING SYSTEM: Madintosh 6.0.5

SOFTWARE: WordPerfect CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/696,793A
FILING DATE: 19910507
CLASSIFICATION: 435
FRIOR APPLICATION DATA:
APPLICATION NUMBER: 32704
FILING DATE:
ATTORNEY/AGENT INPORMATION:
NAME: Kevin R. Kaster
REGISTRATION NUMBER: 32704
REFERENCE/POCKET NUMBER: 32704
REFERENCE (415) 658-5239
INFORMATION FOR SEQ ID NO: 34: SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Nutley
STATE: New Jersey
COUNTRY: U.S.A.
ZIP: 07110-1199
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette, 3.50 inch, 800 Kb storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COMPUTER: Apple Macintosh
OPERATING SYSTEM: Macintosh 6.0.5
SOFTWARE: WordPerfect
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear
MOLECULE TYPE: Other nucleic acid
US-07-696-793A-34
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         159 GAGCGCCCAGTGCCCTGGCC 178
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: NUCLEIC ACID
STRANDEDNESS: single stranded
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Stacey R. Sias, Ph.D.
REGISTRATION NUMBER: 32,630
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             REFERENCE/DOCKET NUMBER: 87
TELECOMMUNICATION INFORMATION:
TELEPHONE: (510) 814-2863
TELEFAX: (510) 814-2977
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/
FILING DATE: 19921117
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
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                                                                        GENERAL INFORMATION:
APPLICANT: GRIffais, R.
APPLICANT: GRIffais, R.
TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragmente TITLE OF INVENTION: Chlamydia pneumoniae genomic sequence and polypeptides, fragmente TITLE OF INVENTION: thereof and uses thereof, in particular for the diagnosis, preve TITLE OF INVENTION: and treatment of infection
FILE REPRENEUE: 9710-003-999
CURRENT APPLICATION NUMBER: US/09/198,452A
CURRENT FILING DATE: 1998-11-24
NUMBER OF SEQ ID NOS: 6849
SEQ ID NOS: 6849
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-10-027-983-90
Sequence 90, Application US/10027983
Sequence 90, Application US/10027983
Sequence 90, Application US/10027983
Sequence 90, Application US/10027983
GENERAL INFORMATION:
APPLICANT: Kenneth W. Dobie
APPLICANT: Mark P. Roach
TITLE OF INVENTION: ANTISENSE MODULATION OF ESTROGEN RECEPTOR ALPHA EXPRESSION
FILE REFERENCE: RTS-0340
CURRENT APPLICATION NUMBER: US/10/027,983
CURRENT FILING DATE: 2001-12-18
NUMBER OF SEQ ID NOS: 98
SEQ ID NO 90
LENGTH: 20
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US-07-696-793A-34/C

i Sequence 34, Application US/07696793A

patent No. 5220004

GENERAL INFORMATION:
 APPLICANT: Saiki, Randall K.
 TITLE OF INVENTION: Typing
 TITLE OF INVENTION: Typing
 NUMBER OF SEQUENCES: 58
 CORRESPONDENCE ADDRESS:
 ADDRESSEE: Cetus Corporation
 STREE: 1400 Faifty-Third Street
 CITY: Emeryville
 STATE: California
 COUNTRY: U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.6%; Score 15.4; DB 1; Length 20; Best Local Similarity 94.1%; Pred. No. 5.9e+02; Matches 16; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; OTHER INFORMATION: Antisense Oligonucleotide
US-10-027-983-90
                  Sequence 6585, Application US/09198452A
Patent No. 6559294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2343 AAGTGCTGGGATTACAG 2359
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           239 TGAAGGAAACTGGGGAG 255
                                                                                                                                                                                                                                                                                                                                                                ; TYPE: DNA
; ORGANISM: Chlamydia pneumoniae
US-09-198-452A-6585
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 AAGTGCTGAGATTACAG 17
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ORGANISM: Artificial Sequence
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  US-09-198-452A-6585/c
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ATTORNEY/AGENT INFORMATION:
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Pred. No. 6.1e+02;
0; Mismatches 3; Indels
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Sequence 15, Application US/07922723A

Batent No. 536004

GENERAL INFORMATION:
APPLICANT: DATE Mineal H. Polymeropoulos
TITLE OF INVENTION: RIVE HIGHLY INFORMATIVE
TITLE OF INVENTION: REPEAT POLYMORPHIC DNA MARKERS
NUMBER OF SEQUENCES:
CORRESPONDENCE ADDRESS:
STREET: Suite 300, 99 Canal Center Plaza
STREET: Suite 300, 99 Canal Center Plaza
STATES: Virginia
                                                                                                                                                                                              Query Match

0.6%; Score 15.2; DB 1;
Best Local Similarity 85.0%; Pred. No. 6.1e+02;
Matches 17; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ZIP: 22314
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: DOS Text File
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/922,723A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FILING DATE:
CLASSIFICATION: 435
ATTORNEY AGENT INFORMATION:
NAME: D.J. Mills
REGISTRATION NUMBER: 34506
REPERBINGE/DOCKET NUMBER: 717081B
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703 684 1111
INFORMATION FOR SEQ ID NO: 35;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 657
US-07-799-828C-35/c
S. Sequence 35, Application US/07799828C
; Patent No. 5378602
; GENERAL INFORMATION:
                                                                                                                             MOLECULE TYPE: Other nucleic acid
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                                                                                                                                                                                                                                                                                    159 GAGCGCCCAGTGCCCTGGCC 178
INFORMATION FOR SEQ ID NO: 34:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
TYPE: NUCLEIC ACID
STRANDEDNESS: single stranded
                                                                                                                                                                                                                                                                                                                   20 GAGTGCCAGTGCCTGTCC 1
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Best Local Similarity 85.0%;
Matches 17; Conservative (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                           TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: Virginia
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                               US-07-922-723A-35/c
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APPLICANT: Drs. Carl R. Wertl and
APPLICANT: Misael H. Polymeropus
TITLE OF INVENTION: TWENT SECREM HIGHLY INFORMATIVE
TITLE OF INVENTION: TWENT SECREM HIGHLY INFORMATION: THE SECREM HIGHLY INFORMATION WOMEN: J. M. MISSIE DOST THE TELECOMMUNICATION HOMEN: J. M. MISSIE DOST THE SECREM HIGHLY INFORMATION: THE SECREM HIGHLY INFORMATION HOMEN: J. M. MISSIE DOST THE SECREM HIGHLY INFORMATION: THE SECREM
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Query Match 0.6%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 6.1e+02; Matches 17; Conservative 0; Mismatches 3; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ORGANISM: Artificial Sequence
FEATURE:
1 OTHER INCOMMATION: Antisense Oligonucleotide
US-09-289-267-165
                                                                                                                                                                           1377 GGCTTTGATGTTCCTGATTG 1396
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ORGANISM: Artificial Sequence
              ORGANISM: Mus musculus
                                                                                                                                                                                                                                                                                           RESULT 661
US-09-289-267-162/c
                                  US-09-104-497-4
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APPLICANT: TOCQUE, Bruno
APPLICANT: TOCQUE, Bruno
APPLICANT: TOCQUE, Bruno
TITLE OF INVENTION: TRANSGENIC ANIMALS OVEREXPRESSING MDM2
FILE REFERENCE: A2716A-US
CURRENT PILING DATE: 1998-06-25
CURRENT FILING DATE: 1998-06-25
EARLIER APPLICATION NUMBER: 60/051,739
EARLIER FILING DATE: 1997-07-03
NUMBER OF SEQ ID NOS: 7
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 3
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: WASTLYK, Bohdan
APPLICANT: WASTLYK, Bohdan
APPLICANT: TOCQUE, Bruno
APPLICANT: ALKGALAF, MOUSSA
TITLE OF INVENTION: TRANSGENIC ANIMALS OVEREXPRESSING MDM2
FILE REFERENCE: A2716A-US
CURRENT APPLICATION NUMBER: US/09/104,497
CURRENT FILING DATE: 1998-06-25
FALLER APPLICATION NUMBER: 60/051,739
RARLIER PILING DATE: 1997-07-03
NUMBER OF SEQ ID NOS: 7
SOFTWARR: PALENTIN Ver. 2.0
SEQ ID NO 4
SEQ ID NO 4
                                                                                                                                                                                                                                                                                                                                     Query Match 0.6%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 6.1e+02; Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match 0.6%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 6.1e+02; Matches 17; Conservative 0; Mismatches 3; Indels
NAME: D.J. Mills
REGISTRATION NUMBER: 34506
REFERENCE/DOCKET NUMBER: 717081C
TELECOMMUNICATION INFORMATION:
TELEPHONE: 703 684 1111
INFORMATION FOR SEQ ID NO: 35:
SEQUENCE CHARACTERISTICS:
LENGTH: 201
                                                                                                                                                                                                                                                                                                                                                                                                                               2096 TTTTGAGACCGAGTCTTGCT 2115
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Patent No. 6028245
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 Acrcrrrrdaagagarccr 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               20 TCTTGAGACAGGGTCTTGCT 1
                                                                                                                                                                                                                                        TOPOLOGY: linear
MOLECULE TYPE: DNA (genomic)
                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: single
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; ORGANISM: Mus musculus
US-09-104-497-3
                                                                                                                                                                                                                                                                                         US-07-952-277A-35
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Sequence 165, Application US/09289267A

Sequence 165, Application US/09289267A

Patent No. 6046320

GENERAL INFORMATION:

APPLICANT: Lex M. Cowsert

TILLO PO INVENTION: ANTISENSE MODULATION OF MDMX EXPRESSION

FILE REFERENCE: RTS-0049

CURRENT APPLICATION NUMBER: US/09/289,267A

CURRENT FILING DATE: 1999-04-04

NUMBER OF SEQ ID NOS: 166

SEQ ID NO 165

LENGTH: 20
Sequence 162, Application US/09289267A

Sequence 162, Application US/09289267A

Patent No. 6046320

GENERAL INFORMATION:

APPLICANT: Brett P. Monia

TITLE OF INVENTION: ANTISENSE MODULATION OF MDMX EXPRESSION

FILE REFERENCE: RTS-0049

CURRENT FILING DATE: 1999-04-04

NUMBER OF SEQ ID NOS: 166

SEQ ID NO 162

LENGTH: 20
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US-09-429-323-4
; Sequence 4, Application US/09429323A
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; OTHER INFORMATION: Description of Artificial Sequence: Probe
US-09-357-740-8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Donna T. Ward
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF RECQL2 EXPRESSION
FILE REFERENCE: RTS-0207
FILE REFERENCE: RTS-0207
CURRENT APPLICATION NUMBER: US/09/798,096
CURRENT FILIG DATE: 2001-03-01
NUMBER OF SEQ ID NOS: 89
                                                                                                                              Length 20;
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                                                                                                                                                                   Indels
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85.0%; Pred. No. 6.1e+02;
tive 0; Mismatches 3;
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Pred. No. 6.1e+02;
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Pred. No. 6.1e+02;
0; Mismatches 3;
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85.0%; Pred. No. v...
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                                                                          OTHER INFORMATION: Antisense Oligonucleotide
                                                                                                                                                                                                    2146 TCTTGGCTCACTGCAAGCTC 2165
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                      TYPE: DNA
ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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ORGANISM: Artificial Sequence
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Best Local Similarity 85.09
Matches 17; Conservative
                                                                                                                                               Best Local Similarity 85.0
Matches 17; Conservative
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Best Local Similarity 85.0
Matches 17; Conservative
                                                                                           US-09-488-856A-71
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US-09-798-096-16
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US-09-357-740-8
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LENGTH: 20
       LENGTH: 20
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                                                           FEATURE:
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CURRENT APPLICATION NUMBER: US/09/488,856A
CURRENT FILING DATE: 2000-01-21
NUMBER OF SEQ ID NOS: 88
SEQ ID NO 71
Patent No. 6140126

Batent No. 640123

GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: Lex M. Cowestr
TITLE OF INVENTION: ANTISENSE MODULATION OF Y-BOX BINDING PROTEIN 1 EXPRESSION FILE REFERENCE: RTS-0092
CURRENT APPLICATION NUMBER: US/09/429,323A
CURRENT FILING DATE: 1999-10-26
NUMBER OF SEQ ID NOS: 89
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APPLICANT: Madeline M. Butler
APPLICANT: Madeline Wyatt
APPLICANT: Jacqueline Wyatt
APPLICANT: Lex M. Cowsert
TITLE OF INVENTION: ANTISENSE MODULATION OF PEPCK-CYTOSOLIC EXPRESSION
TITLE OF INVENTION: ANTISENSE WOOULATION OF PEPCK-CYTOSOLIC EXPRESSION
CURRENT APPLICATION WHORE: US/09/488,671A
CURRENT FILING DATE: 2000-01-21
SEQ ID NO 109
LENGTH: 20
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0.6%; Score 15.2; DB 1;
Best Local Similarity 85.0%; Pred. No. 6.1e+02;
Matches 17; Conservative 0; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0; Mismatches
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Patent No. 6316259
GENERAL INFORMATION:
APPLICANT: Brett P. Monia
APPLICANT: Madeline M. Butler
APPLICANT: Jacqueline Watter
APPLICANT: Jacqueline Watter
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 109, Application US/09488671A Patent No. 6187545 GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                296 GGTGAGGAGCAGCAAATGT 315
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Matches 17; Conservative
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US-09-488-856A-71/c
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                                                                                                                                                                            SEQ ID NO 4
LENGTH: 20
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APPLICANT: Jacquelline Wyatt
TITLE OF INVENTION: ANTISENSE MODULATION OF BH3 INTERACTING DOMAIN DEATH AGONIST
TITLE OF INVENTION: EXPRESSION
FILE PERFRENCE: RTS-0135
CURRENT APPLICATION NUMBER: US/09/657,346A
CURRENT FILING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 174
SEQ ID NO 32
LENGTH: 20
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                                                                                                                                                                                                                      sequence 79, Application US/U9733294A

Patent No. 6492171

GENERAL INPORATION:
APPLICANT: Brett P. Monia
APPLICANT: William Gaade
APPLICANT: William Gaade
APPLICANT: William Gaade
APPLICANT: Susan M. Freder
APPLICANT: Edward V. Wancewicz
TITLE OP INVENTION: ANTISENSE MODULATION OF TERT EXPRESSION
FILE REFERENCE: ISPH-05.7
CURRENT APPLICATION NUMBER: US/09/733,294A
CURRENT APPLICATION NUMBER: 09/572,423
PRIOR FILING DATE: 2000-15-16
NUMBER OF SEQ ID NOS: 108
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.6%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 6.1e+02; Matches 17; Conservative 0; Mismatches 3; Indels
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    Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ; OTHER INFORMATION: Antisense Oligonucleotide US-09-657-346A-32
  0; Mismatches
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US-09-657-346A-33
; Sequence 33, Application US/09657346A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 32, Application US/09657346A, Patent No. 6503754; GENERAL INFORMATION: APPLICANT: Hong Zhang
                                                                                                                                                                                         US-09-733-294A-79/c
; Sequence 79, Application US/09733294A
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                                                 2048 TTTTTTCTTAAATATGTAT 2067
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Matches 17; Conservative
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US-09-657-346A-32
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GENERAL INFORMATION:

APPLICANT: KANAGAMA, TAKAHIRO
APPLICANT: KANAGAMA, TAKAHIRO
APPLICANT: KANAGAMA, TAKAHIRO
APPLICANT: YANADA, KAZUTAKA
APPLICANT: YOKOMAKU, TOYOKAZU
APPLICANT: WONTONION: METHOD FOR DETERMINING A CONCENTRATION OF TARGET NUCLEIC ACID MOI
TITLE OF INVENTION: MUCLEIC ACID PROBES FOR THE METHOD, AND METHOD FOR ANALYZING DAT
TITLE OF INVENTION: THE METHOD
FILE REPRENCE: 19995JUSOXDIV
CURRENT APPLICATION NUMBER: US 09/556,127
PRIOR FILING DATE: 2000-11-29
PRIOR FILING DATE: 1999-04-20
PRIOR FILING DATE: 1999-04-20
NUMBER OF SEQ ID NOS: 70
SOFTWARE: PATENTIN VERSION 3.1
SEQ ID NO 4.2
LENGTH: 20
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                                                                                                                                                      US-09-137-223A-14

Sequence 14, Application US/09137223A

Sequence 14, Application US/09137223A

Patent No. 6420525

GENERAL INFORMATION:

APPLICANT: Yee, David P

APPLICANT: Delsher, Theresa A

TITLE OF INVENTION: TESTIS-SPECIFIC TRANSCRIPTION FACTOR

TITLE REFERENCE: 97-18

CURRENT APPLICATION NUMBER: US/09/137,223A

CURRENT FILING DATE: 1998-08-19

PRIOR PILING DATE: 1997-08-19

NUMBER OF SEQ ID NOS: 16

SEQ ID NO 14

LENGTH: 20
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US-09-137-223A-14
                      2349 TGGGATTACAGGCATGAGCC 2368
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Patent No. 6492121
                                                                     1 TAGGATTACAGGTGTGAGCC 20
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CHER INFORMATION: SYNTHETIC DNA
US-09-725-265-42
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ORGANISM: ARTIFICIAL SEQUENCE
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Best Local Similarity
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US-09-725-265-42/c
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                                                                                                                  Length 20;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 1100 NO. 059513/LN GLEDE KORD, BIGHTH FLOOT CITY: Arlington STATE: Virginia COUNTRY: US
COUNTRY: US
COMPUTER READABLE FORM:
MEDIUM TYPE: Flopy disk
COMPUTER: IBM PC COMPALIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
SOFTWARE: Patentin Release #1.0, Version #1.25 (EPO)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/060,299
FLIING DATE: 15-APR-1998
CLASSIFICATION DATA:
APPLICATION NUMBER: US 60/043,553
FRIUM DATE: 15-ARR-1997
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: US 60/048,740
FILING DATE: 05-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: B.J.SAGGEF
                                                                                                                Query Match 0.6%; Score 15.2; DB 1; Length 2 Best Local Similarity 85.0%; Pred. No. 6.1e+02; Matches 17; Conservative 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
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1100 No. 6545137th Glebe Road, Eighth Floor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Hey, Patricia
APPLICANT: Kawaguchi, Yoshihiko
APPLICANT: Marriman, Tony R
APPLICANT: Merzher, Michael L
TIPLE OF INVENTION: No. 6545137el Receptor
                              ; PEATURE:
** OTHER INFORMATION: Antisense Oligonucleotide
US-09-657-346A-66
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 36,663
REFERENCE/DOCKET NUMBER: 620-35
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2293 AGGATGGTCTCGATCTCCTG 2312
                                                                                                                                                                                                           2285 TGTTAGCCAGGATGGTCTCG 2304
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; Patent No. 6545137
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                                                                                                                                                                                                                                                                                                                                                                                                                                   Todd, John A
Hess, John W
Caskey, Charles T
Cox, Roger D
Gerhold, David
Hammond, Holly
Hey, Patricia
       ORGANISM: Artificial Sequence
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(703)816-4100
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 20 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: nucleic acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
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APPLICANT:
APPLICANT:
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APPLICANT:
APPLICANT:
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APPLICANT: Hong Zhang
APPLICANT: Hong Zhang
APPLICANT: Jacqueline Wyatt
TITLE OF INVENTION: ANTIERSSION
FILE REFERENCE: RTS-0135
CURRENT PELLING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 174
SEQ ID NO 66
LENGTH: 20
Patent No. 6503754
GENERAL INFORMATION:
APPLICANT: Hong Zhang
APPLICANT: Jacqueline Wyatt
TITLE OF INVENTION: ANTIERNSE MODULATION OF BH3 INTERACTING DOMAIN DEATH AGONIST
TITLE OF INVENTION: EXPRESSION
FILE REPERENCE: RTS-0135
CURRENT PILLING NUMBER: US/09/657,346A
CURRENT FILLING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 174
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 49, Application US/09657346A
Patent No. 6503754
GENERAL INFORMATION
GENERAL INFORMATION:
GENERAL INFORMATION: ANTIBENSE
APPLICANT: Jacqueline Wyatt
TITLE OF INVENTION: ANTIBENSE MODULATION OF BH3 INTERACTING DOMAIN DEATH AGONIST
TITLE OF INVENTION: EXPRESSION
FILE REFERENCE: RTS-0135
CURRENT FILING NUMBER: US/09/657,346A
CURRENT FILING DATE: 2000-09-07
NUMBER OF SEQ ID NOS: 174
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Outlier 1, Best Local Similarity 85.0%; Pred. No. 6.1e+02; Matches 17; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FEATURE:
; OTHER INFORMATION: Antisense Oligonucleotide
US-09-657-346A-49
                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION: Antisense Oligonucleotide US-09-657-346A-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           US-09-657-346A-66
; Sequence 66, Application US/09657346A
; Batten No. 6503754
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2146 TCTTGGCTCACTGCAAGCTC 2165
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                          TYPE: DNA ORGANISM: Artificial Seguence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-657-346A-49
                                                                                                                                                                                                                            SEQ ID NO 33
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                         FEATURE:
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1 AGGCTGGTCTCAAACTCCTG 20

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Gaps STATE: Virginia COUNTRY: US ZIP: VA 22201-4714 COMPUTER READABLE FORM: MEDLIUM TYPE: Floppy disk COMPUTER: IBM PC compatible COMPUTER: IBM PC compatible COMPUTER: DATE: IBM PC compatible COMPUTER: DATE: IBM PC compatible SOFTWARE: PATENTIA Release #1.0, Version #1.25 (EPO) CURRENT APPLICATION DATA: APPLICATION DATA: APPLICATION NUMBER: US 60/042, 553 FILING DATE: 15-APR-1997 APPLICATION NUMBER: US 60/048, 740 Gerhold, David
Hammond, Holly
Haw, Patricia
Kawaguchi, Yoshihiko
Merziman, Tony R
Merzer, Michael L
TITLE OF INVENTION. No. 6555654el LDL-Receptor
NUMBER OF SEQUENCES: 455
CORRESPONDENCE ADDRESS:
ADDRESSEE: Nixon and Vanderhye
STREET: 1100 No. 6555654th Glebe Road, Eighth Floor
CITY: Arlington ö Query Match 0.6%; Score 15.2; DB 1; Length 20; Best Local Similarity 85.0%; Pred. No. 6.1e+02; Matches 17; Conservative 0; Mismatches 3; Indels Kawaguchi, Yoshihiko Merriman, Tony R Metzker, Michael L TITLE OF INVENTION: NO. 6555654el LDL-Receptor NUMBER OF SEQUENCES: 455 REGISTRATION NUMBER: 36,663
REPERBYCE POCKET NUMBER: 620-81
TELECOMMULICATION INFORMATION:
TELEPHONE: (703)816-4091
INFORMATION FOR SEQ. ID NO: 78:
SEQUENCE CHARACTERISTICS: ; SEQUENCE DESCRIPTION: SEQ ID NO: 78: US-09-402-923A-78 RESULT 678
US-09-402-923A-240/c
; Sequence 240, Application US/09402923A
; Patent NO. 6555654
; GENERAL INFORMATION:
; APPLICANT: Todd, John A
; Hess, John A
; Caskey, Charles T
Caskey, Charles T 2293 AGGATGGTCTCGATCTCCTG 2312 1 Adderidererezakacieerig 20 Cox, Roger D Gerhold, David Hammond, Holly Hey, Patricia TYPE: nucleic acid STRANDEDNESS: single Roger D TOPOLOGY: linear ð

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Gaps

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GENERAL INPORMATION:
APPLICANT: KURANE, RYUICHIRO
APPLICANT: KANAGAMA, TAKAHIRO
APPLICANT: KANAGATA, YOLCHI
APPLICANT: KANADA, KAZUTAKA
APPLICANT: YOKOMAKU, TOYOKAZU
APPLICANT: YOKOMAKU, TOYOKAZU
APPLICANT: KOYAMA, OSAMU
APPLICANT: FURUSHO, KENTA
TITLE OF INVENTION: METHOD FOR DETERMINING A CONCENTRATION OF TARGET NUCLEIC ACID MOI TITLE OF INVENTION: THE METHOD FOR ANALYZING DATITLE OF INVENTION: THE METHOD
FILE REFERENCE: 0163-0758-0X
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 98, Application US/09953318
Fatent No. 6710174
GENERAL INFORMATION:
APPLICANT: C. Frank Bennett
APPLICANT: C. Frank Bennett
APPLICANT: Andrew T. Watt
TITLE OF INVENTION: ANTISENSE MODULATION OF VASCULAR ENDOTHELIAL GROWTH FACTOR RECEP
TITLE FFERENCE: RTS-0232
TITLE FFERENCE: RTS-0232
CURRENT APPLICATION NUMBER: US/09/953,318
CURRENT FILING DATE: 2001-09-13
NUMBER OF SEQ ID NOS: 154
SEQ ID NO 98
LENGTH: 20
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 20;
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85.0%; Pred. No. 6.1e+02;
tive 0; Mismatches 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/556,127
CURRENT FILING DATE: 2002-06-17
PRIOR APPLICATION NUMBER: 19 1999-111601
PRIOR FILING DATE: 1999-04-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2234 CACCACACCTGGCTAATTTT 2253
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                                                                                                              . Sequence 42, Application US/09556127; Patent No. 6699661
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  20 CTTTTGCCCAGTCTGGAGTG 1
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OTHER INFORMATION: SYNTHETIC DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQ ID NOS: 70
SOFTWARE: Patentin version 3.1
SEQ ID NO 42
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: ARTIFICIAL SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 85.0°
Matches 17; Conservative
                                                                   RESULT 680
US-09-556-127-42/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-556-127-42
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APPLICANT: C. Frank Bennett
APPLICANT: Susan M. Freier
TITLE OF INVENTION: ANTISENSE MODULATION OF SPHINGOSINE-1-PHOSPHATE LYASE EXPRESSION
FILE REFERENCE: RTS-0259
CURRENT APPLICATION NUMBER: US/09/967,669
CURRENT FILING ANTE: 2001-09-28
NUMBER OF SEQ ID NOS: 90
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                                                            CITY: Arington
CITY: Arington
STATE: Virginia
COUNTY: US
ZIE: VA 2201-4714
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25 (EFO)
CURRENT APPLICATION NUMBER: US/09/402,923A
FILING DATE: 14-Feb-2001
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ö
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                        ADDRESSEE: Nixon and Vanderhye
STREET: 1100 No. 6555654th Glebe Road, Eighth Floor
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Length 20;
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0.6%; Score 15.2; DB 1;
Best Local Similarity 85.0%; Pred. No. 6.1e+02;
Matches 17; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.6%; Score 15.2; DB 1; Best Local Similarity 85.0%; Pred. No. 6.1e+02; Matches 17; Conservative 0; Mismatches 3;
                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: PCT/GB98/01102
FILING DATE: 15-APR-1998
APPLICATION NUMBER: US 60/043,553
FILING DATE: 15-APR-1997
APPLICATION NUMBER: US 60/048,740
FILING DATE: 05-JUN-1997
ATTORNEY/AGENT INFORMATION:
NAME: B.J.Sadoff
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Antisense Oligonucleotide US-09-967-669-58
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY: linear
SEQUENCE DESCRIPTION: SEQ ID NO: 240:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 36,663
REFERENCE/DOCKET NUMBER: 620-81
TELECOMMUNICATION INFORMATION:
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Sequence 58, Application US/09967669
Patent No. 6692960
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (703)816-4100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 20 base pairs
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                INFORMATION FOR SEQ ID NO: 240:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ORGANISM: Artificial Sequence PEATURE:
  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-09-402-923A-240
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQ ID NO 58
LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ઠે
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                                                                                                                                                                                                             TYPE: RNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           RESULT 685
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APPLICANT: Kinzler, Kenneth
APPLICANT: Vogelstein, Bert
APPLICANT: Zhou, Wei
TITLE OF INVENTION: Gene Expression Profiles in No. 6333152mal and
TITLE OF INVENTION: Cancer Cells
TITLE OF INVENTION: Cancer Cells
FILE REPERBUCE: 01107.74664
CURRENT APPLICATION NUMBER: US/09/081,646
CURRENT APPLICATION NUMBER: 60/047,352
EARLIER APPLICATION NUMBER: 60/047,352
EARLIER APPLICATION NUMBER: 60/047,352
SOFTWARE: PSEC FOR Windows Version 3.0
SOFTWARE: PSEC FOR Windows Version 3.0
SEC ID NO.3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Nucleic Acid Catalysts with Endonuclease Activity
FILE REFERENCE: MHBMDO.884-CURRENT APPLICATION NUMBER: US/09/479,005A
CURRENT APPLICATION DATE: 2000-01-07
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Patent No. 5194428

Patent No. 5194428

; ZAMECNIK, PAULC.

; TITLE OF INVENTION INHIBITION OF INFLUENZA VIRUS
; REPLICATION BY OLIGONUCLECTIDE PHOSPHOROTHIOATES

; WUMBER OF SEQUENCES: 12

; CURRENT APPLICATION DATE: 30-APR-1990

; PRIOR APPLICATION NUMBER: 160,574

FILING DATE: 26-FEB-1988

; APPLICATION NUMBER: 11,894

FILING DATE: 10-JUL-1987

; APPLICATION NUMBER: 867,231

; FILING DATE: 23-MAY-1986
                                                                                                                                                                                                                                                                                                                                                                                                                                              0.6%; Score 15.2; DB 1; Length 20;
85.0%; Pred. No. 6.1e+02;
tive 0; Mismatches 3; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 15;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 15; DB 1; Lo
Pred. No. 6.9e+02;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 684
US-09-479-005A-139/c
Sequence 139, Application US/09479005A
; Patent No. 6656791
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1111 AACTCTCAGATGAAGATGAT 1130
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-09-081-646-3/c
; Sequence 3. Application US/09081646
; Patent No. 6333152
; GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.6%; Scc
Best Local Similarity 100.0%; Pi
Matches 15; Conservative 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          2350 GGGATTACAGGCATG 2364
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Best Local Similarity 85.0
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                 LENGTH: 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-09-081-646-3
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TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "101858 PROBE; SYNTHETIC
DESCRIPTION: NUCLEOTIDE SEQUENCE COMPLEMENTARY TO NUCLEOTIDES 21-39 IN SEQ ID
DESCRIPTION: NO:8 AND SEQ ID NO:9; S'END FLUORESCEIN LABELED CYTOSINE;"
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                                                                                                                                                                                                                                                                                                                                                                         Gaps
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Fatent. No. 594283
GENERAL INFORMATION:
APPLICANT: Kwok, Pui-Yan
APPLICANT: Chen, Xiangning
TITLE OF INVENTION: Method for Nucleic Acid Analysis Using
TITLE OF INVENTION: Pluorescence Resonance Energy Transfer
NUMBER OF SQUENCES:
ADDRESSEE: Howell & Haferkamp, L.C.
STREET: 7733 Porsyth Boulevard, Suite 1400
CITY: St. Louis
                                                                                                                                                                                                                                                                                                                                                                            ö
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OTHER INFORMATION: /note= "N REPRESENTS S' FLUORESCEIN
OTHER INFORMATION: /nabeled CYTOSINE;"
                                                                                                                                                                                                                                                                                                                           Score 15; DB 1; Length 16; Pred. No. 6.7e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ZIP: 63105-1817

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/767,979
FILING DATE: 17-DEC-1996
CLASSIFICATION: 455
                                                                                                                                                                                                                                                                                                           0.6%; Scot.
100.0%; Pred. No. cot.
... 0; Mismatches
              PRIOR FILING DATE: 1999-11-19
PRIOR APPLICATION NUMBER: US 09/159,274
PRIOR PILING DATE: 1998-09-22
PRIOR FILING DATE: 1997-09-22
NUMBER OF SEQ ID NOS: 1208
SOFTWÄRE: Patentin version 3.0
LENGTH: 16
APPLICATION NUMBER: US 09/444,209
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: HOLLand, Donald R
REGISTRATION NUMBER: 35.197
REPERENCE/DOCKET NUMBER: 96-5.
TELECOMMUNICATION INFORMATION:
TELERAN: 314-727-6192
INFORMATION FOR SEQ ID NO: 10:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
TYRE: nucleic acid
TYRE: nucleic acid
STRANDENESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                     453 TATACTATGAAAGAG 467
                                                                                                                                                                                                                                                                                                                                                                                                                                                                16 TATACTATGAAGAG 2
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FEATURE:
NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                    Query Match 0.6%
Best Local Similarity 100.0
Matches 15, Conservative
                                                                                                                                                                                                                                                      , ORGANISM: Homo sapiens
US-09-479-005A-139
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Query Match
      Matches
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                                                                                                                                                    Gaps
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                                                                                                                                                                                                                                                                                                                               Sequence 10, Application US/09295026
Patent No. 6177249
GENERAL INPORMATION:
APPLICANT: Kwok, Pui-Yan
Chen, Xiangning
TITLE OF INVENTION: Method for Nucleic Acid Analysis Using
Fluorescence Resonance Energy Transfer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CORRESPONDENCE ADDRESS:
ADDRESSE: Howell & Haferkamp, L.C.
STREET: 7733 Forsyth Boulevard, Suite 1400
CITY: St. Louis
STATE: MO
CUNTRY: USA
ZIP: 63105-1817
COMPUTER READBALE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
CORPUTER: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/295,026
FILING DATE: 20-Apr-1999
CLASSIFICATION: cUnknown>
                                                                                                       Query Match 0.6%; Score 15; DB 1; Length 19; Best Local Similarity 100.0%; Pred. No. 6.3e+02; Matches 15; Conservative 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.6%; Score 15; DB 1; Length 19; Best Local Similarity 100.0%; Pred. No. 6.3e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
MOLECULE TYPE: other nucleic acid
DESCRIPTION: /desc = "D18S8 PROBE; SYNTHETIC
HYPOTHETICAL: NO
ANTI-SENSE: NO
FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/767,979
FILING DATE: CURROWAN-
ATTORNEY/AGENT INFORMATION:
NAME: Holland, Donald R
REGISTRATION NUMBER: 35,197
REFERENCE/DOCKET NUMBER: 96-5219
TELEPRONCE: 314-727-5188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: 1 - SEQUENCE DESCRIPTION: SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 314-727-6092
INFORMATION FOR SEQ ID NO: 10:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
                                                                                                                                                                                      2155 ACTGCAAGCTCTGCC 2169
                                                                                                                                                                                                                        2 ACTGCAAGCTCTGCC 16
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES: 34
                       misc_feature
1
; FEATURE:
; NAME/KEY:
; LOCATION:
US-08-767-979-10
                                                                                                                                                                                                                                                                                                            -09-295-026-10
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; NAME/KEY: primer_bind
; LOCATION: 1..20
; OTHER INFORMATION: downstream amplification primer 99-14468 for SEQ 351, in complem
US-09-422-978-8216
                                                                                                                                                                                                                                                          APPLICANT: Cohen, Daniel
APPLICANT: Chem, Daniel
APPLICANT: Chem, Daniel
APPLICANT: Chemseld, Marta
APPLICANT: Chumakov, Ilya
TITLE APPLICANT: Chumakov, Ilya
TITLE REFERENCE: GENSET-020CP1
CURRENT APPLICATION NUMBER: US/09/422,978
CURRENT PILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US 09/298,850
EARLIER APPLICATION NUMBER: US 60/109,732
EARLIER APPLICATION NUMBER: US 60/109,732
EARLIER PILING DATE: 1998-11-23
EARLIER PILING DATE: 1998-11-23
EARLIER FILING DATE: 1998-11-23
EARLIER FILING DATE: 1998-11-30
EARLIER FILING DATE: 1998-11-30
EARLIER FILING DATE: 1998-11-30
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Gaps
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Patent No. 6566135

GENERAL INFORMATION:
APPLICANT: Vickie L. Brown-Driver
APPLICANT: Vickie L. Brown-Driver
APPLICANT: Watt

TITLE OF INVENTION: ANTISENSE MODULATION OF CASPASE 6 EXPRESSION
FILE REPERBNCE: RTS-0187
CURRENT APPLICATION NUMBER: US/09/679,299A

CURRENT FILING DATE: 2000-10-04

NUMBER OF SEQ ID NOS: 164

SEQ ID NO 72

LENGTH: 20
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Indels
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O.6%; Score 15; DB 1; Le
Best Local Similarity 100.0%; Pred. No. 6.2e+02;
Matches 15; Conservative 0; Mismatches 0;
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; OTHER INFORMATION: Antisense Oligonucleotide
US-09-679-299A-72
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
                                                                                                                                                                           US-09-422-978-8216
; Sequence 8216, Application US/09422978
; Patent No. 6537751
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     100.0%; Pre
;
                                          2155 ACTGCAAGCTCTGCC 2169
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2318 GIGATCCGCCCACCT 2332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          829 ATGAATTATCTGGTG 843
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5 GIGATCCGCCCACCT 19
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15; Conservative
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Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: DNA
ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        RESULT 688
US-09-679-299A-72
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQ ID NO 8216
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COMPUTER READABLE FORM:

MEDIUM TYPE: 1910A

COMPUTER READABLE FORM:

MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE

COMPUTER: IBM PC.

COMPUTER: DISKETTE: S.0

CURRATING SYSTEM: PC-DOS

SOFTWARE: WORDPERFECT 5.0

CURRATING SYSTEM: 19930121

CLASSIFICATION NUMBER: US/08/007,997A

FILING DATE: 19930121

CLASSIFICATION NUMBER: 939,855

FILING DATE: SEPTEMBER: 939,855

FILING DATE: SEPTEMBER: 2, 1992

PRIOR APPLICATION NUMBER: PCT/US91/05209

FILING DATE: AUJUST: 14, 1990

ATTORNEY/AGRET INFORMATION:

NAME: Jane Massey Licata

REGISTRATION NUMBER: 32,257

REGISTRATION NUMBER: 33,257

REGISTRATION NUMBER: 33,257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
APPLICANT: Bennett and Mirabelli
TITLE OF INVENTION: Oligonucleotide Modulation
TITLE OF INVENTION: Of Cell Adhesion
NUMBER OF SEQUENCES: 82
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock Washburn Kurtz
ADDRESSEE: Mockiewicz & No. 5591623ris
STREET: One Liberty Place - 46th Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match

0.6%; Score 14.8; DB 1;
Best Local Similarity 88.9%; Pred. No. 6.6e+02;
Matches 16; Conservative 0; Mismatches 2;
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Jane Massey Licata
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0002
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          RESULT 691
US-08-007-997A-4/c
; Sequence 4, Application US/08007997A
; Patent No. 5591623
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2322 TCGGCCCACCTCGGCCTC 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REPERENCE/DOCKET NUMBER: 151S
RELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
TELEPAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18 rccrcccaccrcaccrc 1
                                                                                                                                                                                                                                                  SENCENCE IN TRANSPER NACIONAL STRANDEDNESS: Single TOPOLOGY: Linear ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
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                  Sequence 1171, Application US/09443199C

Patent No. 6670464

GENERAL INFORMATION:
APPLICANT: Leach, Martin
TITLE OF INVENTION: Nucleic Acids Containing Single Nucleotide
TITLE OF INVENTION: Polymorphisms and Methods of Use Thereof
FILE REFERENCE: 15966-534A
CURRENT FILING DATE: 1999-11-16
PRIOR APPLICATION NUMBER: 60/109,024
PRIOR FILING DATE: 1998-11-17
NUMBER OF SEQ ID NOS: 1272
SOFTWARE: Curaden Patent Formatter Version 0.9
SEQ ID NO 1171
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
| LOCATION: (26)...(0)
| OTHER INFORMATION: 1 of 2 allelic variants (1172 is other entry)
| NAME/KEY: misc_feature
| LOCATION: (0)...(0)
| OTHER INFORMATION: Accession number cg43129081
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.6%; Score 15; DB 1; Length 51; Best Local Similarity 61.5%; Pred. No. 4.2e+02; Matches 24; Conservative 0; Mismatches 15; Indele
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER REDABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DOS
SOFTWARE: MORDPERPECT 5.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/063,167A
FILING DATE: 19930517
CLASSIFICATION 514
PRIOR APPLICATION NUMBER: 939.855
FILING DATE: September 2, 1992
PRIOR APPLICATION NUMBER: 939.855
FILING DATE: September 2, 1992
PRIOR APPLICATION NUMBER: 939.855
FILING DATE: Supy 23, 1991
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 567,286
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               567 TTTGGCGTGCCAAGCTTCTCTGTGAAAGAGCACAGGAAA 605
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            US-08-063-167A-4/c

Sequence 4, Application US/08063167A

Fatent No. 5514788

GAPLICANT: Bennett and Mirabelli

TITLE OF INVENTION: Oligonucleotide Modulation

TITLE OF INVENTION: of Cell Adhesion

NUMBER OF SEQUENCES: 85

CORRESPONDENCE ADDRESS:

ADDRESSE: Woodland Falls Corporate Park

STREET: 210 Lake Drive East, Suie 201

CITY: Cherry Hill
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               45 TTTGGGAGGCCAAGGTGGGCGGATCACGAGGTCAGGAGA 7
                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
ORGANISM: Homo sapiens
US-09-443-199C-1171/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CITY: Cherr
STATE: NJ
COUNTRY: US
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FILING DATE: August 14, 1990 PRIOR APPLICATION DATA: APPLICATION NUMBER:
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Pred. No. 6.6e+02;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DOS
SOFTWARE: WORDERFECT 5.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/440,740A
FILING DATE: May 12, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-440-740A-4/c

Sequence 4, Application US/08440740A

Patent No. 5843738

GENERAL INFORMATION:

APPLICANT: Bennett and Mirabelli

TITLE OF INVENTION: Oligonucleotide Modulation

TITLE OF INVENTION: of Cell Adhesion

NUMBER OF SEQUENCES: 85

CORRESPONDENCE ADDRESS:

ADDRESSEE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ISPH-0133
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 063,167

FILING DATE: May 17, 1993

PRIOR APPLICATION DATA:
APPLICATION NUMBER: 969,151

FILING DATE: February 10, 1993

PRIOR APPLICATION NUMBER: 007,997

FILING DATE: January 20, 1993

PRIOR APPLICATION NUMBER: 939,855

FILING DATE: September 2, 1992

PRIOR APPLICATION NUMBER: 939,855

FILING DATE: September 2, 1992

PRIOR APPLICATION NUMBER: 939,855

FILING DATE: AUGUST 11,1990

AUGUST 11,1990
                                                                                                                                                                                                                                                                                                        2322 TCCGCCCACCTCGGCCTC 2339
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REGISTRATION NUMBER: 32,257
REPERENCE/DOCKET NUMBER: 1SF
TELECOMMUNICATION INFORMATION:
TELEPHONE: (609) 779-2400
                                                                                                                                                                                                                                                                                                                                                              18 rccrccaccrcagccrc
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEFAX: (609) 779-8488
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         QUENCH: 18
LENGTH: 18
TYPE: Nucleic Acid
LENGTH: 18
TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                     Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Mariton
STATE: NJ
COUNTRY: USA
                                                                                  ; TOPOLOGY: Line
; ANTI-SENSE: Yes
US-08-007-997A-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-440-740A-4
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0.6%; Score 14.8; DB 1; Length 18;

Query Match

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                                 Gaps
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                                                                                                                                                               RESULT 693
US-08-344-155C-4/C
is Sequence 4, Application US/08344155C
is Patent No. 5883082.
is Patent No. 5883082.
is APPLICANT: Bennett and Stepkowski
it TITLE OF INVENTION: Compositions and Methods for Preventing
it TITLE OF INVENTION: OPPOSITIONS OF SEQUENCES: 99
is CORRESPONDENCE ADDRESS:
is ADDRESSE: Woodland Falls Corporate Park
is STREET: 210 Lake Drive East, Suite 201
CITY: Cherry Hill
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                                 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: NGA

COUNTRY: NGA

ZIP: 08002

COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PS/2
COMPUTER: IBM PS/2
COMPUTER: IBM PS/2
COMPUTER: WORDERFECT 5.0

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/344,155C
FILING DATE: SOPTEMBER: 33, 1994

CLASSIFICATION DATA:
APPLICATION NUMBER: PCT/US91/05209
FILING DATE: SOPTEMBER: PCT/US91/05209
FILING DATE: July 23, 1991
PRIOR APPLICATION NUMBER: 08/063,167
FILING DATE: J/1/93
PRIOR APPLICATION NUMBER: 08/063,167
FILING DATE: J/21/93
PRIOR APPLICATION NUMBER: 08/063,167
FILING DATE: 9/2/92
PRIOR APPLICATION NUMBER: 07/939,855
FILING DATE: B/14/90
APPLICATION NUMBER: 07/939,855
FILING DATE: B/14/90
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: 07/567,286
FILING DATE: B/14/90
ATTORNEY/AGENT INFORMATION:
ANDARY: JANEY: JANEY
88.9%; Pred. No. -
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REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0098
TELECOMMUNICATION INFORMATION:
TELEPHONE: (609) 779-2400
                                                                            2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2322 TCGCCCACCTGGCCTC 2339
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INFORMATION FOR SEQ ID NO: 4
SEQUENCE CHARACTERISTICS:
LENGTH: 18
           Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TOPOLOGY: Linear
ANTI-SENSE: Yes
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-344-155C-4
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; TOPOLOGY: linear
US-08-403-888A-120
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: NJ
COUNTRY: US,
ZIP: 08053
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-982-845B-4/c
                                                                                                                                                                                                                                                                                                                                                                                                                       LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
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Sequence 120, Application US/08403888A

Patent No. 5952490

ERNERAL INFORMATION:

APPLICANT: Hanecak et al.

TITLE OF INVENTION: Oligonucleotides Having A Conserved G4 Core

TITLE OF INVENTION: Sequence

NUMBER OF SEQUENCES: 146

CORRESPONDENCE ADDRESS:

ADDRESSEE: Woodcock Washburn Kurtz Mackiewicz & No. 5952490ris LLP

STREET: One Liberty Place - 46th Floor

COUNTRY: U.S.A.

STATE: PA J. S.A.

COUNTRY: U.S.A.

COUNTRY: U.S.A.

CONPUTER READABLE FORM:

MEDIUM TYPE: 3.5 inch disk, 1.44 Mb

COMPUTER: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.6%; Score 14.8; DB 1; Length 18; 88.9%; Pred. No. 6.6e+02; tive 0; Mismatches 2; Indels
                                                                                                                                                   GENERAL INCORRATION:
APPLICANT: Tam, Robert C.
TITLE OF INVENTION: METHODS AND COMPOSITIONS FOR
TITLE OF INVENTION: REGULATION OF CD28 EXPRESSION
NUMBER OF SEQUENCES: 48
CORRESPONDENCE ADDRESS:
ADDRESSEB: Crockett & Fish
STREET: 3000 S. Augusta Court
CITY: La Habra
STRET: La Habra
STRTE: California
COUNTRY: United States of America
ZIP: 90631
                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Wordberfect 6.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/529,878B
FILING DATE: 13-SEP-1995
CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Fish, Robert D.
REGISTRATION NUMBER: 33,880
REFERENCE/DOCKET NUMBER: 213/003
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECHONE: 714-525-3433
                                                                RESULT 694
US-08-529-878B-33/c
; Sequence 33, Application US/08529878B
; Patent No. 5932556
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18 TCCTCCCACCTCAGCCTC 1
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TOPOLLOGY: unknown
MOLECULE TYPE: DNA (genomic)
US-08-529-878B-33
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX:
INFORMATION FOR SEQ ID NO: 33
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
TYPE: nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity 88.9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
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Matches
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READBLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PS/2
OPERATING SYSTEM: Windows 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/982,845B
FILING DATE: December 2, 1997
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/440,740
FILING DATE: May 12, 1995
PRIOR APPLICATION NUMBER: 063,167
PRIOR APPLICATION NUMBER: 969,151
APPLICATION NUMBER: 939,855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 4, Application US/08982845B
; Patent No. 6015894
GENERAL INFORMATION:
GENERAL INFORMATION:
TITLE OF INVENTION: Oligonuclectide Modulation
TITLE OF INVENTION: of Cell Adhesion
NUMBER OF SEQUENCES: 87
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 66 East Main Street
CITY: MarIton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                  SOFTWARE: Wordberfect 6.1

CURRENT APPLICATION DATA:
PELLING DATE: 12-JUN-1995
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: 07/954,185
FILING DATE: 29-SEP-1992
ATTONNEY AGENT INFORMATION:
NAME: Paul K. Legaard
REGISTRATION NUMBER: 38,534
REFERENCE/DOCKET NUMBER: 151S-1229
TELECOMMUNICATION INFORMATION:
TELEPHONE: 215-568-1100
TELEPHONE: 215-568-1100
TELEPHONE: 215-568-1100
TELEPHONE: 215-568-1100
TELEPHONE: 215-568-1100
TELEPHONE: 215-568-1100
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2322 TCCGCCCACCTCGGCCTC 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       18 TCCTCCCACCTCAGCCTC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: nucleic acid
STRANDEDNESS: single
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Matches 16; Conservative
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US-09-109-273-7
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                                                                                                                                                                                                                                                                                                                                            SEQ ID NO 43
LENGTH: 18
                                                                                                                                   RESULT 698
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APPLICANT: Fernandez-Alnemri, Teresa
APPLICANT: Fernandez-Alnemri, Teresa
TITLE OF INVENTION: USING THE SAME, AND COMPOSITIONS FOR AND METHODS
TITLE OF INVENTION: Of MAKING THE SAME
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
ADDRESSEE: Woodcock, Washburn, Kurtz, Mackiewicz & No. 6037461ris
STREET: One Liberty Place, 46th floor
CITY: Philadelphia
                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                                                           ö
                                                                                                                                                                                                                                                                                                                                                              Score 14.8; DB 1; Length 18;
Pred. No. 6.6e+02;
0; Mismatches 2; Indels
          APPLICATION NUMBER: 567,286
FILING DATE: August 14, 1990
ATTORNEY,AGENT INFORMATION:
NAME: Jane Massey Licata
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0243
TELECOMMUNICATION INFORMATION:
TELEPHONE: (609) 779-2400
TELEPHONE: (609) 779-2400
TELEPHONE: (609) 779-8488
INPORMATION FOR SEQ ID NO: 4:
ELROTH: 18
LENGTH: 18
TYPE: NUCLEIC ACID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SOFTWARE: WordPerfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/859,167
FILING DATE:
CLASSIFICATION: 435
ATTORNEY/AGENI INFORMATION:
NAME: Deluca, Mark
REGISTRATION NUMBER: 33,229
REGISTRATION NUMBER: 33,229
REFERENCE/DOCKET NUMBER: TJU-
TELEPHONE: (215) 568-3100
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence 7, Application US/08859167
Patent No. 6037461
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                             2322 TCCGCCCACCTCGGCCTC 2339
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: WINDOWS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   18 rccrcccaccrcaeccrc 1
                                                                                                                                                                                                                                                                                                                                                                  0.6%;
88.9%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: (215) 568-3439
INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: nucleic acid
STRANDEDNESS: sing
                                                                                                                                                                                                                                                                                    ; TOPOLOGY: Linear
; ANTI-SENSE: Yes
US-08-982-845B-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           , MOLECULE TYPE: DNA
US-08-859-167-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  -08-859-167-7
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Score 14.8; DB 1; Length 18; Pred. No. 6.6e+02;

0.6%;

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Patent No. 6063760
GENERAL INFORMATION:
APPLICANT: Alnemri, Emad S.
APPLICANT: Fernandez-Alnemri, Teresa
TITLE OF INVENTION: FADD-LIKE ANTI-APOPTOTIC MOLECULES, METHODS OF
TITLE OF INVENTION: USING THE SAME, AND COMPOSITIONS FOR AND METHODS
TITLE OF INVENTION: OF MAKING THE SAME
NUMBER OF SEQUENCES: 17
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Woodcock, Washburn, Kurtz, Mackiewicz & No. 6063760ris
STREET: One Liberty Place, 46th floor
CITY: Philadelphia
Gaps
                                                                                                                                                                                        "US-09-339-993-43/c
| Sequence 43, Application US/09339993A
| Sequence 43, Application US/09339993A
| General No. 6640179
| General No. 6640179
| The Part No. 6640179 | The Part No. 6640179 | The Part No. 6640179 | The Part No. 6640179 | The Part No. 6640179 | The Part No. 6640179 | CURRENT APPLICATION NUMBER: US/09/339,993A | CURRENT FILING DATE: 1999-06-25 | NUMBER OF SEQ ID NOS: 47
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Pred. No. 6.6e+02;
0; Mismatches 2; Indels
2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ) OTHER INFORMATION: Antisense Oligonucleotide US-09-339-993-43
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 19103
ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYRE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: WINDOWS
SOFTWARE: WordPerfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/109,273
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DAID.

CLASSIFICATION
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/859,167
FILING DATE:
ATTORNEY/ABENT INFORMATION:
NAME: Deluca, Mark
REGISTRATION NUMBER: 33,229
REGISTRATION NUMBER: TJU-
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
TELECOMMUNICATION:
                                                 2293 AGGATGGTCTCGATCTCC 2310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1147 CTGTGTATCAGGCAGGGG 1164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: DNA ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity 88.9%;
Matches 16; Conservative (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    18 crererareaceaces
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Gaps

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Parent No. 6096722
GENERAL INFORMATION:
APPLICANT: C. Frank Bennett, Christopher Mirabelli,
APPLICANT: Erands Baker
TITLE OF INVENTION: Antisense Modulation of Cell Adhesion
TITLE OF INVENTION: Molecule Expression and Treatment of Cell Adhesion
TITLE OF INVENTION: Molecule-Associated Diseases
NUMBER OF SEQUENCES: 109
CORRESPONDENCE ADDRESS:
                                                                                                                           ö
                                                                   Score 14.8; DB 1; Length 18;
Pred. No. 6.6e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PS/2
OPERATING SYSTEM: PC-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/085,759
FILING DATE: herewith
CLASSIFICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/440,740
FILING DATE: May 12, 1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 063,167
FILING DATE: May 17, 1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 969,151
FILING DATE: February 10, 1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 007,997
FILING DATE: January 20, 1993
PRIOR APPLICATION NUMBER: 939,855
FILING DATE: SEPTEMBER: 2, 1992
PRIOR APPLICATION NUMBER: 8199,855
FILING DATE: SEPTEMBER: 2, 1992
PRIOR APPLICATION NUMBER: 8199,855
FILING DATE: AUGUBET 1, 1990
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: 319,1990
ATTORNEY/AGENT INFORMATION:
APPLICATION NUMBER: 319,1990
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Jane Massey Licata
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: ISPH-0311
TELECOMMUNICATION INFORMATION:
TELEPHONE: (609) 779-2400
                                                                                                                                                                        2322 TCCGCCCACCTCGGCCTC 2339
                                                                                                                                                                                                                                                                                                                     US-09-085-759-4/c
; Sequence 4, Application US/09085759
; Patent No. 6096722
                                                                      Query Match
Best Local Similarity 88.9%;
Matches 16; Conservative
                                                                                                                                                                                                                     18 recreceácercasecere
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; TYPE: Nucleic Acid
; STRANDEDNESS: Single
; TOPOLICGY: Linear
US-09-085-759-4
; ANTI-SENSE:
US-08-991-525B-4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PS/2
OPERATING SYSTEM: Windows 95
SOFTWARE: WORDPERFECT 6.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/991,525B
FILING DATE: December 16, 1997
CLASSIFICATION S14
PRIOR APPLICATION NUMBER: 440,740
FILING DATE: May 12, 1995
PRIOR APPLICATION NUMBER: 663,167
FILING DATE: May 17, 1993
PRIOR APPLICATION NUMBER: 969,151
FILING DATE: February 10, 1993
PRIOR APPLICATION NUMBER: 969,151
FILING DATE: February 21, 1993
PRIOR APPLICATION NUMBER: 969,151
FILING DATE: January 21, 1993
PRIOR APPLICATION NUMBER: 567,286
FILING DATE: September 2, 1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 567,286
FILING DATE: September 2, 1992
PRIOR APPLICATION DATA: ANGUST 14, 1990
ATTORNEY/AGENT INFORMATION:
NUMBER: Jane Massey Licatea
                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-991-525B-4/C

Sequence 4, Application US/08991525B

Sequence 4, Application US/08991525B

Sequence 4, Application US/08991525B

Patent No. 6039811

APPLICANT: Bennett and Mirabelli

TITLE OF INVENTION: Oligonucleotide Modulation

TITLE OF INVENTION: Oligonucleotide Modulation

TITLE OF INVENTION: Of Cell Adhesion

NUMBER OF SEQUENCES: 87

CORRESPONDENCE ADDRESS:

ADDRESSE: Law Offices of Jane Massey Licata

STREET: 66 East Main Street

CITY: Mariton

STATE: NJ

COUNTRY: USA
                                                                                                                                                                                                                                                                                           2293 AGGATGGTCTCGATCTCC 2310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Jane Massey Licata
REGISTRATION NUMBER: 32,257
REFRENCE/DOCKET NUMBER: ISPH-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (856) 810-1515
INFORMATION FOR SEO ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                       LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
TOPOLLGY: linear
MOLECULE TYPE: DNA
                                                                                                                                                                                         Ouery Match
Best Local Similarity 88.99
Matches 16; Conservative
SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TYPE: Nucleic Acid
STRANDEDNESS: Single
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RESULT 703
US-09-276-993-7
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                                               Gaps
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Query Match 0.6%; Score 14.8; DB 1; Length 18; Best Local Similarity 88.9%; Pred. No. 6.6e+02; Matches 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 08053
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: IBM PS/2
OPERATING:SYSTEM: PC-DOS
SOFTWARE: MORDERRECT 5.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/128,496
                                                                                                                                                                                                                                                                                                                 APPLICANT: Bennett and Mirabelli
TITLE OF INVENTION: Oligomucleotide Modulation
TITLE OF INVENTION: Oligomucleotide Modulation
TITLE OF INVENTION: Oligomucleotide Modulation
NUMBER OF SEQUENCES: 85
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Offices of Jane Massey Licata
STREET: 65 East Main Street
CITY: Mailton
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRIOR APPLICATION:
PRIOR APPLICATION:
PRIOR APPLICATION NUMBER: 08/440,740
FILING DATE: PRIOR PELICATION NUMBER: 969,151
FILING DATE: PEDTUARY 10, 1993
FILING DATE: PEDTUARY 10, 1993
FILING APPLICATION NUMBER: 907,997
FILING DATE: JANUARY 20, 1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 939,855
FILING DATE: September 2, 1992
PRIOR APPLICATION NUMBER: 939,855
FILING DATE: September 2, 1992
APPLICATION NUMBER: 14, 1990
ATTORNEY AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Jane Massey Licata
REGISTRATION NUMBER: 32,257
REPERENCE/DOCKET NUMBER: 15PH-
TELECOMMUNICATION INFORMATION:
TELEPAN: (609) 779-2400
TELEPAN: (609) 779-8488
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2322 TCCGCCCACCTCGGCCTC 2339
                                                                                                                                                                                                                              US-09-128-496-4/c
; Sequence 4, Application US/09128496
; Patent No. 6169079
; GENERAL INFORMATION:
                                                                                            2322 TCCGCCCACCTCGGCCTC 2339
                                                                                                                                      18 rccrcccaccrcagccrc 1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 16; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; ANTI-SENSE: Yes
US-09-128-496-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 USA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: N
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Sequence 7, Application US/09276993
| Sequence 7, Application US/09276993
| Patent No. 6207801
| GENERAL INFORMATION:
| APPLICANT: Alnemri, Emad S. APPLICANT: Alnemri, Emad S. APPLICANT: Alnemri, Teresa APPLICANT: Fernandez-Alnemri, Teresa TITLE OF INVENTION: USING THE SAME, AND COMPOSITIONS FOR AND METHODS OF TITLE OF INVENTION: Of MAKING THE SAME NUMBER OF SEQUENCES: 17
| CORRESPONDENCES: 17
| ADDRESSEE: Woodcock, Washburn, Kurtz, Mackiewicz & No. 6207801ris STREET: One Liberty Place, 46th floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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Patent No. 6238863

GENERAL INFORMATION:
APPLICANT: Schumm, James W.
APPLICANT: Bacher, Jeffery W.
TITLE OF INVENTION: MATERIALS AND METHODS FOR
TITLE OF INVENTION: IDENTIFYING AND ANALYZING INTERMEDIATE TANDEM
TITLE OF INVENTION: REPEAT DNA MARKERS
NUMBER OF SEQUENCES: 147
CORRESPONDENCE ADDRESS:
ADDRESSEE: Promega Corporation
STREET: 2800 Woods Hollow Road
COTTY: Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ö
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Pred. No. 6.6e+02;
0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                 CITY: Philadelphia
STATE: PA
COUNTRY: USA
ZIP: 19103
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: WINDOWS
SOFTWARE: WOOTDERfect
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/276,993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/859,167
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY AGENT INFORMATION:
NAME: DeLuca, Mark
REGIESTRATION NUMBER: 33,229
REFRENCE/DOCKET NUMBER: TJU-
TELECOMMUNICATION INFORMATION:
TELEPHONE: (215) 568-3100
INFORMATION FOR SEQ ID NO: 7:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  2293 AGGATGGTCTCGATCTCC 2310
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match.

Best Local Similarity 88.9%;
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TOPOLOGY: linear;
; MOLECULE TYPE: DNA
US-09-276-993-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STATE: Wisconsi
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     RESULT 704
US-09-018-584A-69/c
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.6%; Score 14.8; DB 1; Length 18; Best Local Similarity 88.9%; Pred. No. 6.6e+02; Matches 16; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPOTER READABLE FORM:
MEDIUM TYPE: DISKETTE, 3.5 INCH, 1.44 Mb STORAGE
COMPUTER: 1BM PG. Windows 95
SOFTWARE: WORDPERFECT 6.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Bennett and Mirabelli
TITLE OF INVENTION: Oligonucleotide Modulation
TITLE OF INVENTION: of Cell Adhesion
NUMBER OF SEQUENCES: 95
CORRESPONDENCE ADDRESS:
ADDRESSEE: Law Office of Jane Massey Licata
STREET: 66 East Main Street
CITY: Marlton
STATE: Was
COUNTY: USA
ZIP: 08053
COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb
COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb
COMPUTER: IBM compatible FC
OPERATING SYSTEM: Windows 95
SOFTWARE: Word 97 (DOS text format)
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/018,584A
FILING DATE: 04-Feb-1998
CLASSIFICATION:
NAME: Grady J. Frenchick
REGISTRATION NUMBER: 29,018
REFERENCE/DOCKET NUMBER: 16026.9180
TELECOMMUNICATION INFORMATION:
TELEPHONE: (608) 257-3501
TELEPHONE: (608) 257-3501
INFORMATION FOR SEQ ID NO: 69:
SEQUENCE CHARACTERISTICS:
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SCHWARE: WORDPERECT 6.
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/009,490A
FILING DATE: January 20, 1998
CLASSIFICATION: 514
PROR APPLICATION DATA:
APPLICATION NUMBER: 440,740
FILING DATE: MAY 12, 1995
RIOR APPLICATION DATA:
APPLICATION NUMBER: 063,167
FILING DATE: MAY 17, 1993
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 969,151
FILING DATE: February 10, 1993
RAPLICATION DATA:
APPLICATION NUMBER: 939,855
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-009-490A-4/c
; Sequence 4, Application US/09009490A
; Patent No. 6300491
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2110 CTTGCTCTGTTACCCAGG 2127
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            18 CTTGCTCTCTTGCCCAGG 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: Nucleic Acid
STRANDEDNESS: Single
TOPOLOGY: Linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-018-584A-69
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RESULT '0',
US-09-784-423-69/C
'US-09-784-423-69, Application US/09784423
; Patent No. 6767703
; GENERAL INFORMATION:
; APPLICANT: Schumm, James W.
; Bacher, Jeffery W.
: TITLE OF INVENTION: MATERIALS AND METHODS FOR IDENTIFYING AND ANALYZING INTERMEDIATE TANDEM
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
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Best Local Similarity 88.9%; Pred. No. 6.6e+02;
Matches 16; Conservative 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                Query Match 0.6%; Score 14.8; DB 1; Best Local Similarity 88.9%; Pred. No. 6.6e+02; Matches 16; Conservative 0; Mismatches 2;
FILING DATE: September 2, 1992
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 567,286
FILING DATE: August 14, 1990
ATTORNEY/AGENT INFORMATION:
NAME: Jane MASSEY Licata
REGISTRATION NUMBER: 32,257
REFERENCE/DOCKET NUMBER: 1SPH-0268
TELECOMMINICATION INFORMATION:
TELEFAX: (609) 810-1515
TELEFAX: (609) 810-1454
INFORMATION FOR SEQ ID NO: 4:
SEDUENCE CHARACTERISTICS:
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: DNA
ORGANISM: Artificial Sequence
FRATURE:
NAME/KEY: misc_feature
OTHER INFORMATION: No. 6576751el Sequence
US-09-723-450-7
                                                                                                                                                                                                                                                                                                                                                                                                                   2322 TCGGCCCACCTCGGCCTC 2339
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                                                                                                                                                                                                                                                                                                                                                                                                                                                    18 recreceacereacere 1
                                                                                                                                                                                                                                           TYPE: Nucleic Acid
STRANDENNESS: Single
TOPOLOGY: Linear
JANT-SENSE: Yes
US-09-009-490A-4
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TOPOLOGY: Linear

ANTI-SENSE: Yes
PCT-US93-08101-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Madison
STATE: Wisconsin
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-018-584A-64/c
                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH:
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                             CORRESPONDENCES: 14,

NUMBER OF SEQUENCES: 14,

ADDRESSEE: Promega Corporation
STREET: 2800 Woods Hollow Road
CITY: Madison
STATE: Wadison
STATE: Wisconsin
COUNTRY: U.S.A.

ZIP: 53711-5399
COMPUTER READABLE FORM:
MEDIUM TYBE: Diskette - 3.5 inch, 1.44 Mb
COMPUTER: IBM compatible PC
COMPUTER: IEM Compatible PC
COMPUTER: IS-Feb-2001
CLASSIFICATION NUMBER: US/09/784,423
FILING DATE: 15-Feb-2001
CLASSIFICATION NUMBER: 09/018,584
FILING DATE: 04-Feb-1998
ATTORNEY/AGENT INFORMATION:
NAMBE: Grady J. Frenchick
REGISTRATION NUMBER: 16026,9180
TELEPHONE: (608) 257-3501
TELEPHONE: (608) 257-2275
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COMPUTER READABLE FORM:
COMPUTER READABLE FORM:
COMPUTER: IBM PS/2
COMPUTER: IBM PS/2
COMPUTER: IBM PS/2
CORRATING SYSTEM: PC-DOS
CONTWARE: WORDPERFECT 5.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/08101
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Oligonucleotide Modulation TITLE OF INVENTION: Oligonucleotide Modulation TITLE OF INVENTION: of Cell Adhesion NUMBER OF SEQUENCES: 85
CORRESPONDENCE ADDRESS:
ADDRESSE: Woodland Falls Corporate Park STRET: 210 Lake Drive East, Suie 201
CITY: Cherry Hill STATE: NJ
REPEAT DNA MARKERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: Linear; SEQUENCE DESCRIPTION: SEQ ID NO: 69 US-09-784-423-69
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Sequence 4, Application PC/TUS9308101
GENERAL INFORMATION:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER: 939,855
FILING DATE: September 2, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2110 CTTGCTCTGTTACCCAGG 2127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             INFORMATION FOR SEQ ID NO: 69
SEQUENCE CHARACTERISTICS:
                      SEQUENCES: 147
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                      NUMBER OF
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APPLICATION NUMBER: 107/1091/05209
PRIOR APPLICATION NUMBER: 557,286
PRIOR APPLICATION NUMBER: 557,286
PRIOR APPLICATION NUMBER: 557,286
PRIOR APPLICATION NUMBER: 557,286
PRIOR APPLICATION NUMBER: 12,257
PRIOR APPLICATION NUMBER: 12,257
PRESENCE APPLICATION NUMBER: 13,257
PRESENCE APPLICATION NUMBER: 13,267
PRESENCE APPLICATION NUMBER: 15,267
PRESENCE APPLICATION NUMBER: 15,27
PRESENCE APPRESENCE APPLICATION NUMBER: 15,27
PRESENCE APPLICATIO
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US-09-784-423-64/C
; Sequence 64, Application US/09784423
; Patent No. 6767703
; Patent INFORMATION:
; GENERAL INFORMATION:
; APPLICANT: Schumm, James W.
; TITLE OF INVENTION: MATERIALS AND METHODS FOR
; IDENTIFYING AND ANALYZING INTERMEDIATE TANDEM
REPEAT DNA MARKERS
                                                                                                                    Gaps
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                                                                Query Match 0.6%; Score 14.8; DB 1; Length 19; Best Local Similarity 88.9%; Pred. No. 6.5e+02; Matches 16; Conservative 0; Mismatches 2; Indels
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                                                                                                                                                                                                                                                                                                                                                   | February No. 96 / 391/
| GENERAL INFORMATION:
| APPLICANT: Korneluk, Robert G.
| APPLICANT: LaCasse, Eric
| APPLICANT: Baird, Stephen
| APPLICANT: Bolik, Martin
| APPLICANT: Young, Sean
| TITLE OF INVENTION: Antisense IAP Nucleic Acids and Uses
| TITLE OF INVENTION: Antisense IAP Nucleic Acids and Uses
| TITLE OF INVENTION: Antisense IAP Nucleic Acids and Uses
| TITLE OF INVENTION: Thereof
| FILE REFERENCE: 07891/025001
| CURRENT PILING DATE: 2010-09-28
| NUMBER OF SEQ ID NOS: 231
| SEQ ID NO 128
| LIBUGTH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSE: Promega Corporation
STREET: 2800 Woods Hollow Road
CITY: Madison
STATE: Wisconsin
COUNTRY: U.S.A.
ZIP: 53711-5399
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette - 3.5 inch, 1.44 Mb
COMPUTER: IBM compatible PC
OPERATING SYSTEM: Windows 95
SOFTWARE: Word 97 (DOS text format)
CURRENT APPLICATION DATA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/784,423
FILING DATE: 15-Feb-2001
CLASSIFICATION: «Unknown»
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; OTHER INFORMATION: based on Homo sapiens US-09-672-717-128
                                                                                                                                                                                                                                                                                                                    Sequence 128, Application US/09672717
Patent No. 6673917
                                                                                                                                                          2106 GAGICTTGCTCTGTTACC 2123
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Artificial Sequence
Linear
       ;
US-09-018-584A-64
                                                                                                                                                                                                                                                                                                 US-09-672-717-128
TOPOLOGY:
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Sequence 721-6174
Sequence 721-6174
Sequence 721-6134
GENERAL INFORMATION:
APPLICANT: RObbins, Joan M.
APPLICANT: Tritz, Richard
TITLE OF INVENTION: RIBOZYME THERAPY FOR THE TREATMENT OF PROLIFERATIVE
TITLE OF INVENTION: SKIN AND EYE DISEASES
FILE REFERENCE: 480124-40.7
CURRENT APPLICATION NUMBER: US/09/696,791
CURRENT APPLICATION NUMBER: US/09/696,791
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 2142
LENGTH: 19
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APPLICANT: Giordano, J.Y.
TITLE OF TOWNENTION: Expressed Sequence Tags and Encoded Human Proteins.
Patent No. 6783961
PILE REFERENCE: 59.U52.REG
CURRENT APPLICATION NUMBER: US/09/513,999C
CURRENT FILING DATE: 2000-02-24
PRIOR APPLICATION NUMBER: US 60/122,487
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0.6%; Score 14.8; DB 1; Length 19;
88.9%; Pred. No. 6.5e+02;
tive 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ; OTHER INFORMATION: Cyclin E ribozyme binding site US-09-696-791-2142
               APPLICATION NUMBER: 09/018,584
FILING DATE: 04-Feb-1998
ATTORNEY/AGENT INFORMATION:
NAME: Grady J. Frenchick
REGISTRATION NUMBER: 29,018
REFERENCE/DOCKET NUMBER: 16026.9180
TELECOMMUNICATION INFORMATION:
TELEPAN: (608) 257-3501
TELEPAN: (608) 257-3501
TELEPAN: (608) 257-2275
INFORMATION FOR SEQ ID NO: 64
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: Linear
SEQUENCE DESCRIPTION: SEQ ID NO: 64
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 16016, Application US/09513999C
; Patent No. 6783961
; GENERAL INFORMATION:
; APPLICANT: Dumas Milne Edwards, J.B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2106 GAGTCTTGCTCTGTTACC 2123
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1517 CATTATTATAGCAGCCA 1534
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                                                                                                                                                                                                                                                                                                                             TYPE: Nucleic Acid
STRANDEDNESS: Single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              19 GAGTCTCGCTCTGTTGCC
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.6
Best Local Similarity 88.9
Matches 16; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                           LENGIH: 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       RESULT 713
US-09-513-999C-16016/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-09-696-791-2142
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FILE REFERENCE: MBHB00-884-C
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US-09-531-000-72/c
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                                                                                                                                                                                                                                                                       SEQ ID NO 260
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 15118, Application US/09513999C
| Retent No. 6783961
| GENERAL INFORMATION:
| APPLICANT: Dundert, A. APPLICANT: Duclert, A. APPLICANT: Duclert, A. APPLICANT: Duclert, A. APPLICANT: Duclert, A. TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins. | Patent No. 6783961 | Patent No. 6783961 | FILE REFERENCE: 59. US2.REG | CURRENT APPLICATION NUMBER: US 60/122,487 | PRIOR FILING DATE: 1999-02-24 | PRIOR FILING DATE: 1999-02-26 | NUMBER OF SEQ.ID NOS: 36681
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14 AGGTGGATGGATCACAAGGTCAGGAGTTTGAGACCAGCCTGGCCAACATGCTGAAACCCT 73
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Sequence 260, Application US/09479005A
Sequence 260, Application US/09479005A
Sequence 260, Application US/09479005A
Sequence 260, Application:
Sequence 260, Application:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Nucleic Acid Catalysts with Endonuclease Activity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ö
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0.6%; Score 14.8; DB 1; Length 100;
Best Local Similarity 51.5%; Pred. No. 2.9e+02;
Matches 34; Conservative 0; Mismatches 32; Indels (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 0.6%; Score 14.8; DB 1; Length 81;
Best Local Similarity 67.9%; Pred. No. 3.3e+02;
Matches 19; Conservative 0; Mismatches 9; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   891 GATGAAAGCCTGGCTCTGTGTAATAA 918
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            gargaacccrcrcrcrcracrraaannaa s
                                                                                                                                                                                                                                                           FEATURE:
NAME/KEY: misc_feature
LOCATION: 8
OTHER INFORMATION: n=a, g, c or t
                                                                                                                                                                                                                                     OTHER INFORMATION: n=a, g, c or t
PRIOR FILING DATE: 1999-02-26
NUMBER OF SEQ ID NOS: 36681
SOFTWARE: Patent.pm
SEQ ID NO 16016
LENGTH: 81
                                                                                                                                                                                                                                                                                                                                                                                                                            ) OTHER INFORMATION: r=a or g
US-09-513-999C-16016
                                                                                                                                                                 FEATURE:
NAME/KEY: misc_feature
LOCATION: 7
LOCATION: 7
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; ORGANISM: Homo sapiens
US-09-513-999C-15118
                                                                                                                          TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
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LENGTH: 100
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Sequence 271, Application US/09479005A
Farent No. 6656731
GENERAL INFORMATION
TITLE OF INVENTION: Nucleic Acid Catalysts with Endonuclease Activity
TILLS REFRENCE: MBHB00-884-C
CURRENT APPLICATION NUMBER: US/09/479,005A
CURRENT FILING DATE: 1999-11-19
FRIOR APPLICATION NUMBER: US 09/444,209
FRIOR PILING DATE: 1999-11-19
FRIOR PILING DATE: 1998-09-22
FRIOR FILING DATE: 1998-09-22
FRIOR FILING DATE: 1999-09-22
FRIOR FILING DATE: 1999-09-22
FRIOR FILING DATE: 1999-09-22
FRIOR FILING DATE: 1990-09-22
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1; Indels
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68.8%; Pred. No. 7.2e+02;
tive 4; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 14.4; DB 1;
Pred. No. 7.2e+02;
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APPLICANT: JOHNSON, Marion D.
APPLICANT: FRESCO, Jacques R.
TITLE OF INVENTION: TRIPLEX IN-SITU HYBRIDIZATION
CURRENT APPLICATION NUMBER: US/09/479,005A
CURRENT FILING DATE: 2000-01-07
PRIOR PAPLICATION NUMBER: US 09/44,209
PRIOR PILING DATE: 1999-11-19
PRIOR PILING DATE: 1999-09-22
PRIOR FILING DATE: 1998-09-22
PRIOR FILING DATE: 1990-09-22
NUMBER OF SEQ ID NOS: 1208
SOFTWARE: PATENTIN VETSION 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3; Mismatches
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CURRENT APPLICATION NUMBER: US/09/531,000
CURRENT FILING DATE: 2000-09-08
PRIOR APPLICATION NUMBER: PCT/US98/23765
PRIOR PILING DATE: 1998-11-10
PRIOR APPLICATION NUMBER: 60/064,997
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Best Local Similarity 75.0%;
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Best Local Similarity 68.8
Matches 11; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          12; Conservative
                                                                                                                                                                                                                                                                                                                                                                      ; TYPE: RNA
; ORGANISM: Homo sapiens
US-09-479-005A-260
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: RNA
ORGANISM: Homo sapiens
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TITLE OF INVESTION MYCHATION MYCHAILED.

TITLE REFERENCE: AECOMICA-7

CURRENT PILING DATE: 2001-05-25

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR FILING DATE: 2000-09-27

PRIOR APPLICATION NUMBER: PCT/US01/00666

PRIOR APPLICATION NUMBER: PCT/US01/00667

PRIOR APPLICATION NUMBER: PCT/US01/00667

PRIOR APPLICATION NUMBER: PCT/US01/00664

PRIOR FILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00669

PRIOR APPLICATION NUMBER: PCT/US01/006
                                                                                                                                                                                                                                                           GENERAL INFORMATION:
APPLICANT: GU, Yiahong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANNON, Mark
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 14.4; DB 1; Length 17; Pred. No. 7.1e+02;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                          Dequence 7369, Application US/09866108A Patent No. 6686188
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Patent No. 5496699
GENERAL INFORMATION:
TITLE OF INVENTION: Detection of
TITLE OF INVENTION: Gene Sequences
TITLE OF INVENTION: Fluids
TITLE OF INVENTION: Fluids
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
CORRESPONDENCE ADDRESS:
ADDRESSE: Lahive & Cockfield
STREET: 60 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       819 GAAAATTCAGATGAAT 834
           2 AGAAAATTCAGTTGAA 17
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Best Local Similarity 93.8
Matches 15; Conservative
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APPLICANT: GU, Yizhong

APPLICANT: JI, Yongqang

APPLICANT: PENN, Sharron G.

APPLICANT: RANK, David K.

APPLICANT: RANK, David R.

APPLICANT: RANK, David R.

APPLICANT: SHANNON, Wark

TITLE OF INVENTION: MASSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE

PILE REFERENCE: AEOMICA-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILE AREKENNES: ABONULCA-1
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT FILING DATE: 2001-05-25
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PILING DATE: 2000-09-27
PRIOR PELING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/US01/0066
PRIOR PELING DATE: 2001-01-30
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                                                                                                                                                                                                                                                               FEATURE:
OTHER INFORMATION: Description of Artificial Sequence: Target
OTHER INFORMATION: sequences
US-09-531-000-72
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                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match 0.6%; Score 14.4; DB 1; Best Local Similarity 93.8%; Pred, No. 7.1e+02; Matches 15; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 7367, Application US/09866108A Patent No. 668618B
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1305 GAAGATAAAGGGAAAG 1320
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   1997-11-10
                                                                                                                                                                                         TYPE: DNA ORGANISM: Artificial Sequence
                              NUMBER OF SEQ ID NOS: 77
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 72
LENGTH: 17
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Best Local Similarity 93.8
Matches 15; Conservative
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PRIOR FILING DATE:
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US-09-866-108A-7367
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INFORMATION FOR SEQ ID NO: 6
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
; INFORMATION FOR SEQ ID NO: ; SEQUENCE TCARACTERISTICS: ; LENGTH: 18 base pairs ; TYPE: nucleic acid ; STRANDEDNESS: single ; TOPOLGY: linear US-08-551-275-9
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Best Local Similarity 93.8
Matches 15; Conservative
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US-08-483-746A-6
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  US-08-483-746A-6
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US-09-137-075-1
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US-08-551-275-9
; Sequence 9, Application US/08551275
; Patent No. 5679551
; Patent No. 5679551
; Patent No. 7579571
; TILLE OF INVENTION: Unique Double-Stranded RNAs Associated
; TILLE OF INVENTION: With the Trichomonas Vaginalis Virus
; VORRESPONDENCE ADDRESS:
; ADDRESSEE: ADDRESSEE:
; ADDRESSEE: P.O. Box 4433
; CITY: Houston

CITY: May a sequence of the sequence of 
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COUNTRY: US
ZIP: 77210
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/551,275
FILING DATE: 30-0CT-1995
CLASSIFICATION: 514
ATTONEY/AGERT INFORMATION:
                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy Disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCII Text
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/142,845
                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
CLASSIFICATION: 435
CLASSIFICATION DATA:
APPLICATION NUMBER: 07/874,845
FILING DATE: 27-APR-1992
ATTORNEY/AGENT INFORMATION:
NAME: William C. Geary III
REGISTRATION NUMBER: 31,357
REFERENCE/DOCKET NUMBER: DCI-037
TELEPHONE: (617) 227-7400
TELEPHONE: (617) 227-7401
INFORMATION FOR SEQ ID NO: 5:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Corder, Timothy S.
REGISTRATION NUMBER: 38,414
REFERENCE/DOCKET NUMBER: UTS
TELECOMMUNICATION INFORMATION:
TELEPHONE: (512) 418-3000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       743 AGAGAAACCTTCATCT 758
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: nucleic acid
STRANDEDNESS: single
   Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOLECULE TYPE: DNA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
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TELEX: N/A
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; Patent No. 6020124
; GENERAL INFORMATION:
; TITLE OF INVENTION:
; TITLE OF INVENTION:
; CORRESPONDERS 24
; CORRESPONDERS 24
; CORRESPONDERS 124
; CORRESPONDERS E1
; TREET: 60 State Street, suite 510
; TREET: 60 State Street, suite 510
; TATE: Doston
; TATE: Massachusetts
; COUNTRY: USA
ZIP: 02109-1875
; COMPUTER FORM:
; MEDIUM TYPE: Floppy disk
; COMPUTER: IBM PC compatible
; COMPATIBLE
; COMPUTER: IBM PC COMPATIB
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Query Match 0.6%; Score 14.4; DB 1; Length 18; Best Local Similarity 93.8%; Pred. No. 7e+02; Matches 15; Conservative 0; Mismatches 1; Indels
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CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/483,746A
FILING DATE: 07-JUN-1995
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 08/142,845
FILING DATE: 25-OCT-1993
ATTORNEY/AGENT INFORMATION:
NAME: Jean M. Silveri
REGISTRATION NUMBER: 39,030
REFERENCE/DOCKET NUMBER: DCI-037CNCP
TELECOMMULICATION INFORMATION:
TELECOMMULICATION INFORMATION:
TELECOMMULICATION INFORMATION:
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US-09-696-791-1868/c

US-09-696-791-1868/c

Sequence 1868, Application US/09696791

Patent No. 6770633

GENERAL INFORMATION:

APPLICANT: Robbins, Joan M.

APPLICANT: Richard

TITLE OF INVENTION: RIBOZYME THERAPY FOR THE TREATMENT OF PROLIFERATIVE

TITLE OF INVENTION: SKIN AND EYE DISEASES

TITLE OF INVENTION: US/09/696,791

CURRENT APPLICATION NUMBER: US/09/696,791

CURRENT FILING DATE: 2000-10-25

NUMBER: PatentIN Ver. 2.0

SEQ ID NO 1868

LENGTH: 19
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                                                                                       COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk

COMPUTER: IN PC compatible

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:

PRIOR APPLICATION DATA:

RILING DATE: 16-Jun-2000

PRIOR APPLICATION NUMBER: US 60/032,224

FILING DATE: 02-DEC-1996

APPLICATION NUMBER: US 08/980,872

FILING DATE: 01-DEC-1997

APPLICATION NUMBER: CT/US97/21992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 19;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match 0.6%; Score 14.4; DB 1; Best Local Similarity 93.8%; Pred. No. 6.8e+02; Matches 15; Conservative 0; Mismatches 1;
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Best Local Similarity 93.8%; Pred. No. 6.8e+02;
Matches 15; Conservative 0; Mismatches 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME: Michael S. Tuscan
REGISTRATION NUMBER: 43,210
REFERENCE/DOCKET NUMBER: 36870-5057-US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear

MOLECULE TYPE: other nucleic acid

SEQUENCE DESCRIPTION: SEQ ID NO: 20:
US-09-596-377A-20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION
TELEPHONE: 202 467 7000
TELEFAX: 202 467 7176
INFORMATION FOR SEQ ID NO: 20:
SEQUENCE CHARACTERISTICS:
LENGTH: 19 base pairs
    STREET: 1800 M St., NW
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2322 TCCGCCCACCTCGGCC 2337
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1217 TTGGAAATGCACTTCA 1232
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: nucleic acid
STRANDEDNESS: single
                             CITY: Washington
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEATURE:
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US-09-596-377A-20/c

Sequence 20, Application US/09596377A

Sequence 20, Application US/09596377A

Sequence 20, Application US/09596377A

Patent No. 6602850

SENERAL INFORMATION:

APPLICANT: MAGAININ PHARMACEUTICALS, INC.

TITLE OF INVENTION: Biological Variability of Asthma

Associated Factors Useful in Treating and Diagnosing

Atopic Allergies Including Asthma and Related Disorders
                                                                                                                                                                                                                                                                                                                                                                                                                                                               ·;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            sequence 1, Application US/09717793
patent No. 6610486
GENERAL INFORMATION:
APPLICANT: Dahlhauser, Paul
TITLE OF INVENTION: Methods of Screening Nucleic Acids for TITLE OF INVENTION: Mucleotide Variations
FILE REPRENCE: 07036.00002
CURRENT APPLICATION NUMBER: US/09/717,793
CURRENT PILING DATE: 2000-11-20
PRIOR APPLICATION NUMBER: PCT/US99/19007
PRIOR PILING DATE: 1999-08-18
PRIOR FILING DATE: 1998-08-20
NUMBER OF SEQ ID NOS: 2
SOFTWARE: FastSEQ for Windows Version 3.0
SEQ ID NO 1
LENGTH: 18
                                                                                                                                                                                                                                                                                                                                                                                                           0.6%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 7e+02; tive 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0.6%; Score 14.4; DB 1; Length 18; 93.8%; Pred. No. 7e+02; live 0; Mismatches 1; Indels
              TITLE OF INVENTION: Methods of Screening Nucleic Acids for TITLE OF INVENTION: Nucleotide Variations FILE REPERENCE: 07036.001.

CURRENT APPLICATION NUMBER: US/09/137,075

CURRENT APPLING DATE: 1998-08-20

NUMBER OF SEQ ID NOS: 2

SEQ ID NOS: 2

SEQ ID NO 1

LENGTH: 18
                                                                                                                                                                                                                                                                                                     ; NAME/KEY: Oligonucleotide, Single-Stranded
; OTHER INFORMATION: Amplification Primer
US-09-137-075-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NAME/KEY: Oligonucleotide, Single-Stranded; OTHER INFORMATION: Amplification Primer US-09-717-793-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         743 AGAGAAACCTTCATCT 758
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
                                                                                                                                                                                                                                    TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 AGAGAAACCTTTATCT 17
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nes 15; Conservative
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APPLICANT:
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Matches
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TITLE OF INVENTION: Mailelic markers for use in constructing a high density...
FILE REFERENCE: GENSET.020CP1
CURRENT APPLICATION NUMBER: US/09/422,978
CURRENT FILING DATE: 1999-10-20
EARLIER APPLICATION NUMBER: US 60/109,732
EARLIER PILING DATE: 1999-11-23
EARLIER PILING DATE: 1999-11-36
EARLIER PILING DATE: 1999-11-36
EARLIER PILING DATE: 1999-11-36
EARLIER PILING DATE: 1999-11-37
EARLIER APPLICATION NUMBER: US 60/082,614
EARLIER PILING DATE: 1999-121
NUMBER OF SEQ ID NOS: 11796
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: de la Monte, Suzanne
PEDLICANT: Wande, Jack R.
TITLE OF INVENTION: Neural Thread Protein Gene Expression and Detection
TITLE OF INVENTION: of Alzheimer's Disease
NUMBER OF SEQUENCES: 121
CORRESPONDENCE ADDRESS:
ADDRESSE: Sterre, Kessler, Goldstein & Pox P.L.L.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.6%; Score 14.2; DB 1; Length 47; 59.5%; Pred. No. 4.7e+02; tive 1; Mismatches 14; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CONTENTED A LOURENS.
STREET: 1100 New York Avenue, Suite 600
CITY: Washington
STATE: D.C.
COUNTRY: U.S.A.
ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/454,557C
FILING DATE: 30-MAY-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOCATION: 24
CTHER INFORMATION: 99-21516-293 : polymorphic base G or T US-09-422-978-2999
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               785 CTCATCTAGAAGGAGGAGTATTAGTGAGACAGAAGAA 821
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               43 cacecciaraarceraecamirregeaecereaeca 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; Sequence 91, Application US/08454557C; Patent No. 5830670; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELECOMMUNICATION INFORMATION:
TELEPHONE: (202) 371-2600
TELEPAX: (202) 371-2540
INFORMATION FOR SEQ ID NO: 91:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: Ludwig, Steven R.
REGISTRATION NUMBER: 36,203
REFERENCE/DOCKET NUMBER: 06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22; Conservative
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STRANDEDNESS: both
TOPOLOGY: both
                                                                                                                                                                                                                                                                                                                                                                     TYPE: DNA
ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
Matches 22; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: allele
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                                                                                                                                                                                               GENERAL INFORMATION:
APPLICANT: Robbins, Joan M.
APPLICANT: TAILE.
TITLE OF INVENTION: SKIN AND EYE DISEASES
TITLE OF INVENTION: SKIN AND EYE DISEASES
FILE REFERENCE: 480124.407
CURRENT APPLICANT: 2000-10-25
NUMBER OF SEQ ID NOS: 4523
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 3180, Application US/09696791

Batent No. 6770633

GENERAL INFORMATION: Joan M.
APPLICANT: Robbins, Joan M.
APPLICANT: Tritz, Richard

TITLE OF INVENTION: RIBOZYME THERAPY FOR THE TREATMENT OF PROLIFERATIVE

TITLE OF INVENTION: SKIN AND EYE DISEASES
FILE REFERENCE: 480124.407

CURRENT APPLICATION NUMBER: US/09/696,791

CURRENT APPLICATION NUMBER: 2000-10-25

NUMBER OF SEQ ID NOS: 4523

SOFTWARE: Patentin Ver. 2.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match 0.6%; Score 14.4; DB 1; Length 19; Best Local Similarity 93.8%; Pred. No. 6.8e+02; Matches 15; Conservative 0; Mismatches 1; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PEATURE:
; OTHER INFORMATION: Cyclin D1 ribozyme binding site
US-09-696-791-1870
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Cyclin Al ribozyme binding site US-09-696-791-3180
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-422-978-2999/C
; Sequence 2999, Application US/09422978
; Patent No. 6537751
; GENERAL INFORMATION:
; APPLICANT: Cohen, Daniel
; APPLICANT: Blumenfeld, Marta
                                                                                                                                                       Sequence 1870, Application US/09696791
Patent No. 6770633
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1128 GATGAGGTATATCAAG 1143
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19 TTGGAATGAACTTCA 4
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.6
Best Local Similarity 93.8
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       -09-696-791-3180/c
                                                                                                                                  -09-696-791-1870/c
                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQ ID NO 1870
LENGTH: 19
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FILING DATE
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US-08-450-673C-91/c
Sequence 91, Application US/08450673C
Sequence 91, Application US/08450673C
Sequence 91, Application US/08450673C
SENERAL INFORMATION:
APPLICANT: Wands, Jack R.
APPLICANT: Wands, Jack R.
TITLE OF INVENTION: of Alzheimer's Disease
NUMBER OF SEQUENCES: 121
NUMBER OF SEQUENCES: 121
ADDRESSEE: Sterne, Kessler, Goldstein & Fox P.L.L.C.
STREET: 1100 New York Avenue, Suite 600
                                                                                                                                                                                                                   US-08-340-426D-91/c

| Sequence 91, Application US/08340426D
| Patent No. 2946634
| Patent No. 2946634
| GENERAL INFORMATION:
| APPLICANT: de la Monte, Suzanne
| APPLICANT: Wands, Jack R.
| TITLE OF INVENTION: Neural Thread Protein Gene Expression and Detection
| TITLE OF INVENTION: of Alzheimer's Disease
| NUMBER OF SEQUENCES: 121
| CORRESPONDENCE ADDRESS:
                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
                                                                         2277 TTTCACCGTGTTAGCCAGGATGGTCTCGATCTCCTGACCTCGTGATCCGCC 2327
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Pred. No. 3.3e+02;
0; Mismatches 23; Indels
                         0; Mismatches 23; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ADDRESSEE: Sterne, Kessler, Goldstein & Fox P.L.L.C. STREET: 1100 New York Avenue, Suite 600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SIREET: 1100 New YORK AVENUE, SUITE 500
CITY: Washington
STATE: D.C.
COUNTRY: U.S.A.
ZIP: 20005-3934
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: Ploppy disk
COMPUTER: Ploppy disk
COMPUTER: PROPENTION DATA:
COMPUTER: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/340,426D
FILING DATE: 14-NOV-1994
CLASSIFICATION NUMBER: 35,203
FILING DATE: 14-NOV-1994
CLASSIFICATION NUMBER: 36,203
REGISTRATION NUMBER: 36,203
REGISTRATION NUMBER: 36,203
REGISTRATION NUMBER: 36,203
REGISTRATION NUMBER: 36,203
RELEPHONE: (202) 371-2600
INPORMATION FOR SEQ ID NO: 91:
SEQUENCE CHARACTERISTICS:
LEMEGTH: 84 base pairs
LUMBER: A base pairs
  54.9%; Pred. No. 3.3e+02;
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Best Local Similarity 54.9
Matches 28; Conservative
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Best Local Similarity 54.9
Matches 28; Conservative
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STRANDEDNESS:
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US-08-340-426D-91
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TIPL SOURCE READABLE TORN:

MEDIUM TYPE: RIPPE: PLOSPY disk

MEDIUM TYPE: RIPPE: PLOSPY disk

COMPUTE: IRM PC compatible

MACHINE SHARE THE PLOSPY SHARE

TOMESTY AND THE TOWN THE
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Matches
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                                                                                                                                                       0; Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                            2277 TITCACCGIGITAGCCAGGATGGTCTCGATCTCCTGACCTCGTGATCCGCC 2327
                                                                                                                                                                                                                                63 TTTGGGAGGCTGAGCCGGGCGGATCACGAGGTCAGGAGTTCGACACCAGCC 13
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Pred. No. 3.1e+02;
0; Mismatches 28; Indels
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Pred. No. 3.3e+02;
0; Mismatches 8; Indels
                                                                                                        Query Match 0.6%; Score 14.2; DB 1; Length 84; Best Local Similarity 54.9%; Pred. No. 3.3e+02; Matches 28; Conservative 0; Mismatches 23; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 13007, Application US/09621976;
Patent No. 6639063;
GENERAL INFORMATION:
APPLICANT: Dumas Milne Edwards, J.B.
APPLICANT: Obsert, S.
APPLICANT: Giordano, J.Y.
TITLE OF INVENTION: ESTS and Encoded Human Proteins.;
FILE REFERENCE: GENSET.054RR2;
CURRENT APPLICATION NUMBER: US/09/621,976;
CURRENT FILING DATE: 2000-07-21
NUMBER OF SEQ ID NOS: 19335;
SOFTWARE: PATENT.PM
                                                                                                                                                                                                                                                                                                                  US-09-621-976-12767/c

Sequence 12767, Application US/09621976

Patent No. 6639063

GENERAL INFORMATION:

APPLICANT: Dumas Milne Edwards, J.B.

APPLICANT: JODert, S.

APPLICANT: Glordano, J.Y.

TITLE OF INVENTION: ESTS and Encoded Human Proteins.

FILE REFERENCE: GENSET.054PR2

CURRENT FILING DATE: 2000-07-21

NUMBER OF EQ ID NOS: 19335

SOFTWARE: Patent.pm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         890 TGATGAAAGCCTGGCTCTGTGTGTAAT 916
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Best Local Similarity 52.5°
Matches 31, Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match 0.6
Best Local Similarity 70.4
Matches 19; Conservative
TYPE: nucleic acid
STRANDEDNESS: both
TOPOLOGY: both
PCT-US95-17111A-91
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      , ORGANISM: Homo sapiens
US-09-621-976-12767
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LENGTH: 85
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LENGTH: 94
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RESULT 736

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Sequence 413, Application US/09827998

APPLICANT: Shannon, Mark

TITLE OF INVENTION: NOVEL ISOFORMS OF HUMAN PREGNANCY-ASSOCIATED PROTEIN E

FILE REFERENCE: MDhMORF-8

CURRENT PILING DATE: 2001-04-06

PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR PAPLICATION NUMBER: US 60/207,456

PRIOR PAPLICATION NUMBER: US 60/236,359

PRIOR FILING DATE: 2000-05-26

PRIOR FILING DATE: 2000-05-26

PRIOR FILING DATE: 2000-09-27

NUMBER OF SEQ ID NOS: 1881

SOFTWARE: Aeomica Sequence Listing Engine

PATOR APPLICATION NUMBER: US 60/236,359

PRIOR FILING DATE: 2000-09-27

NUMBER OF SEQ ID NOS: 1881

SEQUENCE ASSOCIATED PROTEIN PR
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Patent No. 6656700
GENERAL INFORMATION:
APPLICANT: Shannon, Mark
TITLE OF INVENTION: NOVEL ISOFORMS OF HUMAN PREGNANCY-ASSOCIATED PROTEIN E
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                           APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
TITLE OF INVENTION: Nucleic Acid Catalysts with Endonuclease Activity
FILE REPERENCE: MBH800-884-C
CURRENT APPLICATION NUMBER: US/09/479,005A
CURRENT FILING DATE: 1900-01-07
PRIOR APPLICATION NUMBER: US 09/444,209
PRIOR APPLICATION NUMBER: US 09/159,274
PRIOR FILING DATE: 1998-09-22
PRIOR FILING DATE: 1998-09-22
NUMBER OF SEQ ID NOS: 1208
SOSTWARE: PATENTIN VERSION 3.0
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100.0%; Pred. No. 7.48+02;
iive 0; Mismatches 0; Indels
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Pred. No. 7.6e+02;
4; Mismatches 0; Indels
Sequence 254, Application US/09479005A Patent No. 6656731 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match 0.6%;
Best Local Similarity 71.4%;
Matches 10; Conservative
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ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: RNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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US-09-827-998-414/c
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US-09-104-497-1/c
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APPLICANT: GLASTON:
APPLICANT: GLASTON:
APPLICANT: Shannon, Mark
TITLE OF INVENTION: NOVEL ISOFORMS OF HUMAN PREGNANCY-ASSOCIATED PROTEIN E
FILE REFERENCE: MDAMORF-8
CURRENT APPLICATION WUMBER: US/09/827,998
CURRENT FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,36
PRIOR PILING DATE: 2000-05-26
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR FILING DATE: 2000-05-1891
SOFTWARE: Acomica Sequence Listing Engine
PACENT NO 415
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Batent No. 6656700

GENERAL INFORMATION:
APPLICANT: Gu, Yizhong
APPLICANT: Shannon, Mark
TITLE OF INVENTION: NOVEL ISOFORMS OF HUMAN PREGNANCY-ASSOCIATED PROTEIN E
FILE REFERENCE: MDHAORF-9
CURRENT APPLICATION NUMBER: US/09/827,998
CURRENT FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PILING DATE: 2000-05-26
PRIOR PILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-09-27
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Pred. No. 7.4e+02;
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             CURRENT APPLICATION NUMBER: US/09/827,998
CURRENT FILING DATE: 2001-04-06
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR FILING DATE: 2000-05-26
PRIOR FILING DATE: 2000-09-27
PRIOR FILING DATE: 2000-09-27
NUMBER OF SEQ ID NOS: 1881
SOFTWARE: Acomica Sequence Listing Engine
PACOR TO 6556700
SEQ ID NO 6556700
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; Sequence 415, Application US/09827998
; Patent No. 6656700
                                                                                                                                                                                                                                                                                                                                                        Query Match 0.6%; Sc
Best Local Similarity 100.0%; P
Matches 14; Conservative 0;
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Best Local Similarity 100.
Matches 14; Conservative
FILE REFERENCE: MDhMORF-8
                                                                                                                                                                                                                                                                                      ; ORGANISM: Homo sapiens
US-09-827-998-414
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Homo sapiens
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                                                                                                                                                                                                                                                                      TYPE: DNA
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JOHERAN INCOMENTALION:
JAPELICANT: Sherr, Charles J
APPLICANT: Quelle, Dawn E
JAPELICANT: Quelle, Dawn E
JAPELICANT: Weber, Jason D.
APPLICANT: Roussel, Martine F.
APPLICANT: Roussel, Martine F.
APPLICANT: Prederique, Zindy
JITLE ROPERENCE: 1340-1-023 CIP 1
CURRENT APPLICATION NUMBER: US/09/480,718
CURRENT APPLICATION NUMBER: 09/129,855
EARLIER FILING DATE: 1998-08-06
NUMBER OF SEQ ID NOS: 48
SOFTWARE: Patentin Ver. 2.0
SOFTWARE: Patentin Ver. 2.0
LENGTH: 30
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ORGANISM: Artificial Sequence
FRATURE:
OTHER INFORMATION: Description of Artificial Sequence: Primer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 1, Application US/09104497

Patent No. 6028245

GENERAL INFORMATION:
APPLICANT: WASYLIX' Bondan
APPLICANT: TOCQUE, Bruno
ITILE OF INVENTION: TRANSGENIC ANIMALS OVEREXPRESSING MDM2
FILE REFERENCE: AZIAGAUS
CURRENT APPLICATION WHERE: US/09/104,497

CURRENT APPLICATION NUMBER: US/09/104,497

CURRENT APPLICATION NUMBER: 06/051,739

EARLIER APPLICATION NUMBER: 60/051,739

SARLIER PILING DATE: 1997-07-03

NUMBER OF SEQ ID NOS: 7

SOFTWARE: Patentin Ver: 2.0

SEQ ID NO: 1
                                                                                                                                                                                                                                                                    Length 17;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  5; Indels
                                                                                                                                                                                                                                                                    0.6%; Score 14; DB 1; L. 100.0%; Pred. No. 7.4e+02;
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Pred. No. 6.6e+02;
0; Mismatches 5;
                                                                                                                                                                                                                                                                                                                 Mismatches
, NUMBER OF SEQ ID NOS: 1881
; SOFTWARE: Acomica Sequence Listing Engine
; Patent No. 6656700
; SEQ ID NO 416
; IENGTH: 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-09-480-718-35/c; Sequence 35, Application US/09480718; Patent No. 6407062; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                         100.08;
                                                                                                                                                                                                                                                                                                                                                             2356 ACAGGCATGAGCCA 2369
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match 0.6%;
Best Local Similarity 77.3%;
Matches 17; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                     14 AcAGGCATGAGCCA 1
                                                                                                                                                                                                                                                                                       Best Local Similarity 100.
Matches 14; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: DNA ORGANISM: Mus musculus
                                                                                                                                                                          TYPE: DNA ORGANISM: Homo sapiens
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US-07-990-300-3
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US-09-113-999C-15415/C

Sequence 15415, Application US/09513999C

Patent No. 6783961

GENERAL INFORMATION:

APPLICANT: Dumas Milne Edwards, J.B.

APPLICANT: Duclert, A.

APPLICANT: Glordano, J.Y.

TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.

Patent No. 6783961

CURRENT APPLICATION NUMBER: US/09/513,999C

CURRENT APPLICATION NUMBER: US/09/513,999C

CURRENT APPLICATION NUMBER: US 60/122,487

PRIOR FILING DATE: 1999-02-26

NUMBER OF SEQ ID NOS: 36681

SOFTWARE: Patent.pm

SEQ ID NO 15415

LENTH: 92
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                                                                                                                                                                                                                                                                                                        # PACENT NO. 0.03201.
# GENERAL INFORMATION:
# GENERAL INFORMATION:
# APPLICANT: Dunas Milne Edwards, J.B.
# APPLICANT: Duclert, A.
# APPLICANT: Glordano, J.Y.
# TITLE OF INVENTION: Expressed Sequence Tags and Encoded Human Proteins.
# PATENT NO. 6783961
# FILER REFREENCE: 59.US2.REG
# CURRENT APPLICATION NUMBER: US/09/513,999C
# CURRENT APPLICATION NUMBER: US 60/122,487
# PRIOR FILING DATE: 1999-02-26
# PRIOR FILING DATE: 1999-02-26
# NUMBER OF SEQ ID NOS: 36681
# SOFTWARE: Patent.pm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                            Length 30;
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                                                                                      10; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     66 AGGTCAGGAGTTCAAGACCAGCCTGACCAACATGGTGA 29
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60.5%; Pred. No. 3.2e+02;
ative 0; Mismatches 15;
                                          Query Match 0.6%; Score 14; DB 1; Best Local Similarity 66.7%; Pred. No. 5.9e+02; Matches 20; Conservative 0; Mismatches 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0.6%; Score 14; DB 1; 66.7%; Pred. No. 3.9e+02; iive 0; Mismatches 10
                                                                                                                              1638 GGTCGACCTAAAATGGTTGCATTGTCCAT 1667
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  888 TTTGATGAAAGCCTGGCTCTGTGTAATA 917
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                                                                                                                                                                      30 Géragacacadeacarerregrarrecacar 1
                                                                                                                                                                                                                                                                       Sequence 15107, Application US/09513999C Patent No. 6783961
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 0.6'
Best Local Similarity 66.7'
Matches 20; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; TYPE: DNA
; ORGANISM: Homo sapiens
US-09-513-999C-15415
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                          US-09-513-999C-15107
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQ ID NO 15107
LENGTH: 68
US-09-480-718-35
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Matches
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Sequence 13. Application US/07990100

Sequence 13. Application US/07990100

Sequence 13. Application US/07990100

Sequence 14. Setting the Companies of Day Contaminants by PCR MUNICARY TOWNSTRON Beterion of Day Contaminants by PCR MUNICARY USA STATES NO. Cambridge Day Companies of Sequence Companies Sequence Companies Sequence Companies of Sequence Companies Companies Companies Sequence Companies Companie
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Gaps
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0.6%; Score 13.8; DB 1; Length 17;
Best Local Similarity 88.2%; Pred. No. 7.6e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 2164, Application US/08584040

| Sequence 2164, Application US/08584040
| Patent No. 6346396
| Patent No. 6346396
| Patent Pavco, Pamela
| APPLICANT: Pavco, Pamela
| APPLICANT: Stinchcomb, Dan T. APPLICANT: Stinchcomb, Dan T. APPLICANT: Stinchcomb, Dan T. APPLICANT: Bacobedo, Jaime
| TITLE OF INVENTION: TREATMENT OF DISEASES OR TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS | TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS | TITLE OF INVENTION: GROWTH FACTOR | TITLE OF INVENTION: G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Los Angeles
COUNTY: Los Angeles
COUNTY: U.SA..
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 1.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFRCATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: OCCOber 26, 1995
ATTONNEY, AGENT INFORMATION:
NAME: WALDBLYG, Richard J.
REGISTRATION NUMBER: 32,327
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOPTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIFICATION 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 60/005,974
FILING DATE: Octobar 26, 1995
ATTORNEY, AGENT INPORMATION:
NAME: WARDING, RICHARD 32,327
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   in TELEK: 67-3510
in FORMATION FOR SEQ ID NO: 2163:
sequence CHARACTERISTICS:
LENGTH: 17 base pairs
rype: nucleic acid
stranbeness: single
rype: TOPOLOGY: linear
US-08-584-040-2163
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1316 GAAAGATAAAGGGGAAA 1332
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TELEX: 67
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Sequence 2163, Application US/08584040

Patent No. 6346308;

PATENT NO. 6346308;

APPLICANT: Pavco, Pamela
APPLICANT: Bossignon, James
APPLICANT: Stinchoomb, Dan T.
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Gailte 4700
CITY: LOS Angeles
STATE: California
COUNTRY: U.S.A.
           CORRESPONDENCE ADDRESSES

ADDRESSES: Lyon & Lyon
STREET: 633 West Fifth Street
CITY: Los Angeles
STRATE: California
COUNTRY: U.S.A.
ZIP: 90071-206
COWPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 81000mpatible
COMPUTER READABLE FORM:
MEDIUM TYPE: 81000mpatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTAMER: PREASEQ for Windows 2.0
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/985,162
FILING DATE: 04 December 1997
CLASSIFICATION NUMBER: 60/036,476
FILING DATE: 31 JANUARY 1997
ATTORNEY/AGENT INFORMATION:
NAME: WADDLEY RICHARD J.
NEGISTRATION NUMBER: 32,327
TELECOMMUNICATION INFORMATION:
NAME: WADDLEY RICHARD J.
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 955-0440
TELEPHONE: (213) 955-0440
INFORMATION FOR SEQ ID NO: 799:
SEQUENCE CHARACTERISTICS:
TENGTH: 17 Dase pairs
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         408 TIGCITITGAAGITAIT 424
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
       CORRESPONDENCE ADDRESS
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RESULT 751
US-09-371-772B-708/c
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0.6%; Score 13.8; DB 1; Length 17;
Best Local Similarity 88.2%; Pred. No. 7.6e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INCOMMATION:
APPLICANT: Pavco, Pamela
APPLICANT: Beroco, Dames
APPLICANT: Brinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Stinchcomb, Dan T.
APPLICANT: Becobedo, Jaime
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: GROWTH FACTOR
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: 633 West Fifth Street
STREET: Swite 4700
CITY: LOS Angeles
STREET: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FILING DATE: January 11, 1996
CLASSIPICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION TOWNER: 06/005,974
FILING DATE: OCCODER 26, 1995
ATTOMNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFERENCE/DOCKET NUMBER: 218/064
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
REFERENCE/DOCKET NUMBER: 218/064
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 2164:
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDENNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 2548, Application US/08584040 Patent No. 6346398
                                                                                                                                                                                                                                                                                                                                                                                                                    1315 GGAAAGATAAAGGGGAA 1331
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TELEFX: (7-3510
INFORMATION FOR SEQ ID NO: 254
SEQUENCE CHARACTERISTICS:
LENGTH: 17 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                      US-08-584-040-2164
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APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Scobbedo, Jaime
TITLE OF INVENTION: Method and Reagent for the Treatment of Diseases or Conditions RC
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vascular Endothelial Growth Factor Receptor
TITLE OF INVENTION NUMBER: US /09/371,772B
CURRENT FILING DATE: 1999-08-10
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR APPLICATION NUMBER: US 08/584,040
PRIOR PILING DATE: 1996-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: PatentIn version 3.0
SEQ ID NO 1072
LENGTH: 17
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APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Stinchcomb, Dan
APPLICANT: Secobed, Jaime
APPLICANT: Becobed, Jaime
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jaim
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jii
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: 
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0.6%; Score 13.8; DB 1; Length 17;
Best Local Similarity 11.8%; Pred. No. 7.6e+02;
Matches 2; Conservative 13; Mismatches 2; Indels
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                                                                                            APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; Sequence 1306, Application US/09371772B
; Patent No. 6566127
; GENERAL INFORMATION:
; APPLICANT: Ribozyme Pharmaceuticals, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Sequence 6729, Application US/09371772B
; Patent No. 6566127
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 UACUUUUUUUUUUUUUU 17
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-09-371-772B-1072
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             Patent No. 6566127

GENERAL INFORMATION:
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Ribozyme Pharmaceuticals, Inc.
APPLICANT: Bavozyme Pharmaceuticals, Inc.
APPLICANT: APPLICATION NUMBER: US 60/005,974
PRIOR PILING DATE: 1996-10-06
PRIOR APPLICATION NUMBER: US 60/005,974
PRIOR FILING DATE: 1996-10-06
PRIOR PRIOR FILING DATE: 1996-10-06
PRIOR PRIOR PLING DATE: 1996-10-06
PRIOR PRIOR PLING DATE: 1996-10-06
PRIOR PLING DATE: 1996-10-06
PRIOR PRIOR PLING DATE: 1996-10-06

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match 0.6%; Score 13.8; DB 1; Length 17; Best Local Similarity 88.2%; Pred. No. 7.6e+02; Matches 15; Conservative 0; Mismatches 2; Indels
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; Sequence 709, Application US/09371772B
; Patent No. 6566127
Sequence 708, Application US/09371772B
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TYPE: RNA
CORGANISM: Homo sapiens
US-09-371-772B-708
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; ORGANISM: Homo sapiens
US-09-371-7728-709
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 753
US-09-371-772B-1072
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Sequence 2551, Application US/09866108A
Patent No. 6686189
GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: David R.
APPLICANT: PRN, David R.
APPLICANT: BRANK, David R.
APPLICANT: SHANNON MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEONICA-7
CURRENT PAPLICATION NUMBER: US/09/866,108A
APPLICANT: SHANNON MARK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEONICA-7
CURRENT PAPLICATION NUMBER: US/02/866,108A
CURRENT PAPLICATION NUMBER: US/02/02/03/66
PRIOR PILING DATE: 2000-10-40
PRIOR PLILNG DATE: 2000-10-40
PRIOR PLILNG DATE: 2001-01-04
PRIOR PLILNG DATE: 2001-01-04
PRIOR PLILNG DATE: 2001-01-04
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PRIOR PLILNG DATE: 2001-01-30
PRIOR PLING DATE: 2001-01-30
PRIOR 
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88.2%; Pred. No. 7.6e+02;
tive 0; Mismatches 2; Indels
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0.6%; Score 13.8; DB 1;
Best Local Similarity 35.3%; Pred. No. 7.6e+02;
Matches 6; Conservative 9; Mismatches 2;
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| TELEPHONE: (213) 489-1600 | TELEPAX: (213) 955-0440 | TELEX: 67-3510 | 955-0440 | TELEX: 67-3510 | 955-0440 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 956-04710 | 95
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1 UUGCUUUUAAAGUAAUU 17
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US-09-866-108A-2551
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hes 15; Conserva
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                                                                        APPLICANT: Pavco, Pam
APPLICANT: McSwiggen, Jim
APPLICANT: McSwiggen, Jim
APPLICANT: Barbon, Jim
APPLICANT: Stinchcoun, Dan
APPLICANT: Stinchcoun, Dan
APPLICANT: Stinchcoun, Dan
APPLICANT: Escobedo, Jaime
TITLE OF INVENTION: Levels of Vacuular Endothelial Growth Factor Receptor
TITLE OF INVENTION: Levels of Vacuular Endothelial Growth Factor Receptor
CURRENT FILING DATE: 1996-01.08
PRIOR FILING DATE: 1995-10-26
PRIOR FILING DATE: 1995-01-08
NUMBER OF SEQ ID NOS: 14225
SOFTWARE: Parentin version 3.0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT: ARLTAT, Saghir
APPLICANT: Fell, Patricia
APPLICANT: Fell, Patricia
APPLICANT: McSwiggen, James
TITLE OF INVENTION: ENZYMATIC NUCLEIC ACID TREATMENT
TITLE OF INVENTION: OF DISEASES OR CONDITIONS RELATED
TITLE OF INVENTION: TO LEVELS OF EPIDERMAL GROWTH
TITLE OF INVENTION: PACTOR RECEPTORS
NUMBER OF SEQUENCES: 1877
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: FastSEQ for Windows 2.0
CURRENT APPLICATION NUMBER: US/09/401,063
                                        APPLICANT: Ribozyme Pharmaceuticals, Inc. APPLICANT: Pavco, Pam
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME: Warburg, Richard J.
REGISTRATION NUMBER: 32,327
REFRENCE/DOCKET NUMBER: 230/107
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 799, Application US/09401063
Patent No. 6623962
GENERAL INFORMATION:
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FILING DATE: 04 December 1997
APPLICATION NUMBER: 60/036,476
FILING DATE: 31 January 1997
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    948 AGTGAATCTACAGGGAC 964
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PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ) ORGANISM: Homo sapiens
US-09-371-772B-6729
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LENGTH: 17
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APPLICANT: HANZEL, DAVIA R.
APPLICANT: CHEN, Wensheng
APPLICANT: SHANKON, MAK
TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
CURRENT PELICATION NUMBER: US 60/207, 456
PRIOR PELICATION NUMBER: US 60/207, 456
PRIOR PELICATION NUMBER: US 60/207, 456
PRIOR PELICATION NUMBER: US 60/236, 359
PRIOR PELICATION NUMBER: US 60/236, 359
PRIOR PELICATION NUMBER: PCT/USO1/00666
PRIOR PELING DATE: 2000-09-27
PRIOR PELING DATE: 2001-01-30
PRIOR PEL
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APPLICANT: GU, Yizhong

APPLICANT: GU, Yonggang

APPLICANT: HANZEL, David K.

APPLICANT: RANK, David R.

APPLICANT: RANK, David R.

APPLICANT: SHANNON, Wark

TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE

PILE REFERENCE: ADOMCA-7
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/09/866,108A CURRENT FILING DATE: 2001-05-25
Sequence 6340, Application US/09866108A
Patent No. 6686188
GENERAL INFORMATION:
APPLICANT: UI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 7364, Application US/09866108A Patent No. 6686188
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PRECENT NO. 6686188

GENERAL INFORMATION:
APPLICANT: GU, Yizhong
APPLICANT: GU, Yizhong
APPLICANT: GU, Yizhong
APPLICANT: BENN, Barron G.
APPLICANT: FENN, Barron G.
APPLICANT: HANZEL, David R.
APPLICANT: HANZEL, David R.
APPLICANT: HANZEL, David R.
APPLICANT: GHEN, Wensheng
APPLICANT: GHEN, Wensheng
APPLICANT: GHEN, Wensheng
APPLICANT: SHANNON, Mark
TITILE OF INVENTION: WROSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PRILING DATE: 2000-10-04
PRIOR FILING DATE: 2000-10-04
PRIOR PRILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: PCT/USO1/00666
PRIOR APPLICATION NUMBER: PCT/USO1/00667
                                                                                                                                                            PRIOR FILING DATE: 2000-10-04
PRIOR PLILING DATE: 2000-09-27
PRIOR APPLICATION NUMBER: US 60/236,359
PRIOR PLILING DATE: 2000-09-27
PRIOR PLILING DATE: 2001-01-30
PRIOR PLING DATE: 2001-01-30
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Pred. No. 7.6e+02;
0; Mismatches 2; Indels
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PRIOR APPLICATION NUMBER: PCT/US01/00669
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 2001-01-30
APPLICATION NUMBER: PCT/US01/00665
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APPLICATION NUMBER: PCT/US01/00668
US 60/207,456
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                                                LING DATE: 2000-05-26
PLICATION NUMBER: GB 24263.6
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Best Local Similarity 88.2%;
Matches 15; Conservative
                                                                                                                                        FILING DATE: 2000-10-04
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TYPE: DNA
ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                -09-866-108A-7365
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                                                                                                                                                                          Gaps
                    Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
Patent No. 6686188
SEQ ID NO 7365
LENGTH: 17
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Aeomica Sequence Listing Engine
Patent No. 6686188
SEQ ID NO 7366
LENGTH: 17
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0.6%; Score 13.8; DB 1; Length 17;
Best Local Similarity 88.2%; Pred. No. 7.6e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                              0.6%; Score 13.8; DB 1; Length 17;
88.2%; Pred. No. 7.6e+02;
tive 0; Mismatches 2; Indels
PRIOR APPLICATION NUMBER: PCT/US01/00663 PRIOR FILING DATE: 2001-01-30
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                                                                                                                                                                                                                1 AGCAGAAAATTCAGTTG 17
                                                                                                                                            Query Match
Best Local Similarity 88.27
Matches 15; Conservative
                                                                                                TYPE: DNA ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; ORGANISM: Homo sapiens
US-09-866-108A-7366
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APPLICANT: SHANNON, MARK

TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEOMICA-7
CURRENT APPLICATION NUMBER: US/09/866,108A
CURRENT APPLICATION NUMBER: US 60/207,456
PRIOR APPLICATION NUMBER: US 60/207,456
PRIOR PILING DATE: 2000-05-26
PRIOR PILING DATE: 2000-10-46
PRIOR FILING DATE: 2000-10-10
PRIOR PILING DATE: 2000-10-130
PRIOR PILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00668
PRIOR PILING DATE: 2001-01-30
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US-09-866-100A-9742/c
i Sequence 9742, Application US/09866108A
i Batent No. 6686188
i GENERAL INFORMATION:
i APPLICANT: UI, Yonggang
i APPLICANT: PENN, Sharron G.
i APPLICANT: PENN, Sharron G.
i APPLICANT: PENN, David K.
i APPLICANT: CHEN, Wensheng
i APPLICANT: CHEN, Wensheng
i TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
i FILE REFERENCE: AEOMICA-7
i CURRENT APPLICATION NUMBER: US/09/866,108A
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NUMBER OF SEQ ID NOS: 15755
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0.6%; Score 13.8; DB 1; Length 17;
Best Local Similarity 88.2%; Pred. No. 7.6e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Aeomica Sequence Listing Engine
Patent No. 6686188
SEQ ID NO 7370
; Sequence 7370, Application US/09866108A; Patent No. 6686188
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                820 AAAATTCAGATGAATTA 836
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                                                                 GENERAL INFORMATION:
APPLICANT GV. Yizhong
APPLICANT: JI. Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: PANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wenbheng
                                                                                                                                                                                                                                                                                                                              SHANNON, Mark
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816 GAAGAAATTCAGATGA 832 | ||||||||||||||

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PRIOR FILING DATE: 2001-01-30
PRIOR APPLICATION NUMBER: PCT/US01/00663
PRIOR FILING DATE: 2001-01-30
Remaining Prior Application data removed - See File Wrapper or PALM.
NUMBER OF SEQ ID NOS: 15755
SOFTWARE: Acomica Sequence Listing Engine
Patent No. 6686188
SEQ ID NO 9743
LENGTH: 17
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Sequence 3, Application PC/TUS9311582

GENERAL INFORMATION:
APPLICANT: Letwin Bruce
TITLE OF INVENTION: Detection of DNA Contaminants by PCR
NUMBER OF SEQUENCES: 3
CORRESPONDENCE ADDRESS:
ADDRESSEE: Legal Affairs, Genetics Institute, Inc.
STREET: 87 Cambridge STREET: MA
COUNTRY: MA
COUNTRY: MA
COMPUTRY: USA
STATE: MA
COMPUTRY: CAMBRIDE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 17;
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APPLICANT: John Landers
APPLICANT: John Landers
APPLICANT: John Landers
APPLICANT: David Houseman
APPLICANT: David Houseman
APPLICANT: Alain Charest
TITLE OF INVENTION: Methods and Products Related to
TITLE OF INVENTION: Methods and DNA Analysis
FILE REFERENCE: M0656/7045(HCL/MAT)
CURRENT APPLICANTION NUMBER: US/09/404,912
CURRENT FILING DATE: 1999-09-24
PRIOR APPLICATION NUMBER: US 60/101,757
PRIOR APPLICATION NUMBER: PCT/US99/22283
PRIOR PELING DATE: 1999-09-24
NUMBER OF SEQ ID NOS: 691
SOFTWARE: PastSEQ for Windows Version 3.0
LENGTH: 17
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Best Local Similarity 88.2%; Pred. No. 7.6e+02;
Matches 15; Conservative 0; Mismatches 2;
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0.6%; Score 13.8; DB 1;
Best Local Similarity 88.2%; Pred. No. 7.6e+02;
Matches 15; Conservative 0; Mismatches 2;
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Patent No. 6703228
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2153 TCACTGCAAGCTCTGCC 2169
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ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-866-108A-9743
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          TYPE: DNA
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TITLE OF INVENTION: MYOSIN-LIKE GENE EXPRESSED IN HUMAN HEART AND MUSCLE
FILE REFERENCE: AEGOMICA-7

FULE REFERENCE: AEGOMICA-7

CURRENT PILING DATE: 2001-05-25

PRIOR PILING DATE: 2000-05-26

PRIOR PILING DATE: 2000-05-26

PRIOR PILING DATE: 2000-05-26

PRIOR APPLICATION NUMBER: US 60/236,359

PRIOR PILING DATE: 2000-09-27

PRIOR APPLICATION NUMBER: PCT/US01/00666

PRIOR PILING DATE: 2001-09-27

PRIOR PILING DATE: 2001-01-30

PRIOR APPLICATION NUMBER: PCT/US01/00665
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CURRENT FILING DATE: 2001-05-25

PRIOR PRIOR APPLICATION NUMBER: US 60/207,456

PRIOR PILING DATE: 2000-05-26

PRIOR FILING DATE: 2000-10-04

PRIOR FILING DATE: 2000-10-04

PRIOR PILING DATE: 2000-10-27

PRIOR PILING DATE: 2000-10-27

PRIOR PILING DATE: 2001-01-30

PRIOR PILING DATE: 2001-01-30
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88.2%; Pred. No. 7.6e+02;
iive 0; Mismatches 2; Indels
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APPLICANT: GU, Yizhong
APPLICANT: JI, Yonggang
APPLICANT: PENN, Sharron G.
APPLICANT: HANZEL, David K.
APPLICANT: RANK, David R.
APPLICANT: CHEN, Wensheng
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity 88.2
Matches 15; Conservative
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US-09-866-108A-9742
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Gaps

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APPLICANT: Harley, Calvin B.
APPLICANT: Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ADDRESSEE: Townsend and Townsend and Crew LLP STREET: Two Embarcadero Center, Eighth Floor CITY: San Francisco STATE: California COUNTE: USA ZIP: 94111-834 CONFUTER READABLE FORM: MEDIUM TYPE: Floppy disk COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                      Cech, Thomas R.
Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
Morin, Gregg B.
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Sequence 26, Application US/0933964;
Parent No. 6025198;
GENERAL INFORMATION:
APPLICANT: C. Frank Bennett;
APPLICANT: Lex M. Coweert;
TITLE OF INVENTION: ANTISENSE MODULATION OF SHIP-2 EXPRESSION;
FILE REPERENCE: RTS-0065;
CURRENT APPLICATION NUMBER: US/09/339,964;
CURRENT FILING DATE: 1999-06-25;
NUMBER OF SEQ ID NOS: 47;
SEQ ID NO 26

LENGTH: 18
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0.6%; Score 13.8; DB 1; Length 17;
Best Local Similarity 88.2%; Pred. No. 7.6e+02;
Matches 15; Conservative 0; Mismatches 2; Indels
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: PCT/US93/11582
FILING DATE:
CLASSIFICATION:
ATTORNEY/AGENT INFORMATION:
NAME: Lazar, Steven R.
REGISTRATION NUMBER: 32,618
REFERENCE/DOCKET NUMBER: 5214X-PCT
TELEPROCHOCKET NUMBER: 5214X-PCT
TELEPROCHOCKET NUMBER: 5214X-PCT
TELEPROCHOCKET NUMBER: 5214X-PCT
TELEPROCHOCKET NUMBER: 5214X-PCT
TELEPROCHONICATION INFORMATION:
TELEPROCHOCKET NUMBER: 5214X-PCT
TELEPROCHOCKET NUMBER: 5214X-PCT
TELEPROCHOCKET NUMBER: 5214X-PCT
TELEPROCHOCKET NUMBER: 5214X-PCT
TELEPROCHOCKET NUMBER: 5314X-PCT
TELEPROCHOCHOCKET NUMBER: 5314X-PCT
TELEPROCHOCKET NUMBER: 5314X-PCT
TELEPROCHOCKET NUMBER: 53
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ORGANISM: Chinese Hamster
TISSUE TYPE: Chinese Hamster Ovary
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US-08-974-549A-393
; Sequence 393, Application US/08974549A
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ORIGINAL SOURCE:
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CLONE: consensus seq.
POSITION IN GENOME:
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PCT-US93-11582-3
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Pavco, Pamela
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                                                                                                                                                                  Length 18;
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CORRESPONDENCE ADDRESS:
ADDRESSEE: SWABEY OGILVY RENAULT
STREET: Suite 1600, 1981 McGill College Avenue
                                                                                                                                                          Query Match

0.6%; Score 13.8; DB 1;
Best Local Similarity 88.2%; Pred. No. 7.5e+02;
Matches 15; Conservative 0; Mismatches 2;
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STATE: QC
COUNTY: Action of the county of t
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APPLICANT: TANG, JianQing
APPLICANT: RELANCON, Serge B.
TITLE OF INVENTION: A NOVEL STR MARKER SYSTEM
TITLE OF INVENTION: FOR DNA FINGERPRINTING
; LOCATION: 1..18
; OTHER INFORMATION: /note= "TCP1.17 primer"
US-08-974-549A-393
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APPLICATION NUMBER: US/09/084,120
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; Sequence 4459, Application US/08584040
; Patent No. 6346398
; GENERAL INFORMATION:
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CLASSIFICATION:
PRIOR APPLICATION:
APPLICATION NUMBER:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME:
COTE, France
REGISTRATION NUMBER: 1325:
TELECOMMUNICATION INFORMATION:
TELEPHONE: 514 288-9399
                                                                                                                                                                                                                                                                                                                         2279 TCACCGTGTTAGCCAGG 2295
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; MOLECULE TYPE: oligonucleotide
US-09-084-120-7
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INFORMATION FOR SEQ ID NO: 7.
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
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hes 15, Conservative
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US-09-084-120-7/c
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Matches
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APPLICANT: Gildea, Brian D.
APPLICANT: Gildea, Brian D.
APPLICANT: Gildea, Brian D.
APPLICANT: Gildea, Brian D.
TITLE OF INVENTION: Methods, Kits and Compositions Pertaining To PNA
TITLE OF INVENTION: Molecular Beacons
FILE REPERBUES: BP9702US-CP1
CURRENT APPLICATION UNDER: US/09/179,298
CURRENT FILING DATE: 1998-10-27
BARLIER APPLICATION UNDER: 08/958,532
EARLIER FILING DATE: 1997-10-27
NUMBER OF SEQ ID NOS: 13
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                   APPLICANT: StinchComb, Dan T.
APPLICANT: Bacobedo, Jaime
TITLE OF INVENTION: METHOD AND REAGENT FOR THE
TITLE OF INVENTION: TREATMENT OF DISEASES OR
TITLE OF INVENTION: CONDITIONS RELATED TO LEVELS
TITLE OF INVENTION: OF VASCULAR ENDOTHELIAL
TITLE OF INVENTION: GROWTH FACTOR
NUMBER OF SEQUENCES: 8502
CORRESPONDENCE ADDRESS:
ADDRESSEE: LYON & LYON
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 6; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                      CITY: LOS ANGELES
CITY: LOS ANGELES
STATE: California
COUNTRY: U.S.A.
ZIP: 90071-2066
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: BX CORAGE
COMPUTER: BM COMPATIBLE
COMPUTER: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/584,040
FLIANG DATE: JANUARY 11, 1996
CLASSIFICATION DATA:
APPLICATION NUMBER: 60/005,974
FLING DATE: October 26, 1995
ATTORNEY/AGENT INFORMATION:
NAME: Warburg, Richard J.
REFERENCE/DOCKET NUMBER: 31,327
REFERENCE/DOCKET NUMBER: 218/064
TELEFA: 67-3510
SEQUENCE CHARACTERISTICS:
FUNNAMINO FOR REQ ID NO: 4459:
FUNNAMINO FOR REQ ID NO: 4459:
                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence B, Application US/09179298; Patent No. 6355421; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1385 TGTTCCTGATTGTAAAA 1401
McSwiggen, James
Stinchcomb, Dan T.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 UGUUCCUGAUGGUAACA 17
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Best Local Similarity 52.9%;
Matches 9; Conservative
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STRANDEDNESS: single
TOPOLOGY: linear
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FEATURE:
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APPLICANT: COULT, James M.
APPLICANT: Gildea, Brian D.
APPLICANT: Gildea, Brian D.
APPLICANT: Gildea, Brian D.
APPLICANT: Hydiga-Nielben, Jens J.
TITLE OF INVENTION: Methods, Kits and Compositions Pertaining To PNA
TITLE OF INVENTION: Molecular Beacons
FILE REPERBENCE: BP9702US-CP1
CURRENT APPLICATION NUMBER: US/09/179,298
CURRENT FILING DATE: 1998-10-27
EARLIER APPLICATION NUMBER: 08/958,532
EARLIER APPLICATION NUMBER: 08/958,532
SOFTWARE: Patentin Ver. 2.0
SOFTWARE: Patentin Ver. 2.0
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
                                                                                                                                                                                                                                                                                                                                                                    OTHER INFORMATION: Description of Artificial Sequence:SYNTHETIC PROBE OTHER INFORMATION: OR TARGET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                          0.6%; Score 13.8; DB 1; Length 18;
88.2%; Pred. No. 7.5e+02;
tive 0; Mismatches 2; Indels
                                                                                 FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)
OTHER INFORMATION: 5' Fluorescein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 8, Application US/09179298
Patent No. 6355421
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: 5' Fluorescein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                477 TATCTTGGCCAGTATAT 493
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    2 TATATTGGCCAATATAT 18
                        LENGTH: 18
TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ORGANISM: Artificial Sequence
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                                                                                                                                                                                                                                                                                                                            OTHER INFORMATION: 3' Dabcyl
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                    FEATURE:
NAME/KEY: misc_feature
LOCATION: (9). (10)
OTHER INFORMATION: spacer
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OTHER INFORMATION: spacer
                                                                                                                                                                                                                                                          FEATURE:
NAME/KEY: misc_feature
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SEQ ID NO 8
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Sequence 9, Application US/09179298
; Parent No. 6335421
; GREKRAL INFORMATION:
; APPLICANT: Coull, James M.
; APPLICANT: Glidea, Brian D.
; APPLICANT: Hyldig-Nielsen, Jens J.
; TITLE OF INVENTION: Methods, Kits and Compositions Pertaining To PNA
; TITLE OF INVENTION: Methods, Kits and Compositions Pertaining To PNA
; TITLE OF INVENTION: Molecular Beacons
; TITLE OF INVENTION: Molecular Beacons
; CURRENT FILING DATE: 1998-10-27
; EARLIER APPLICATION NUMBER: 08/958,532
; SARLIER PILING DATE: 1997-10-27
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: Patentin Ver. 2.0
; SEQ ID NO 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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OTHER INFORMATION: Description of Artificial Sequence:SYNTHETIC PROBE
OTHER INFORMATION: OR TARGET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; OTHER INFORMATION: Description of Artificial Sequence:SYNTHETIC PROBE ; OTHER INFORMATION: OR TARGET US-09-179-298-9
                                                                                                                                                                                                APPLICANT: Coull, James M.
APPLICANT: Coull, James M.
APPLICANT: Gildea, Brian D.
APPLICANT: Hyldig-Inclaem, Jens J.
TITLE OF INVENTION: Methods, Kits and Compositions Pertaining To PNA
TITLE OF INVENTION: Molecular Beacons
FILE REFERRNCE: BP9702US-CP1
CURRENT APPLICATION NUMBER: US/09/179, 298
CURRENT FILING DATE: 1999-10-27
EARLIER APPLICATION NUMBER: 08/958,532
EARLIER FILING DATE: 1997-10-27
NUMBER OF SEQ ID NOS: 13
SOFTWARE: PETENTIN Ver: 2.0
SEQ ID NO
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88.2%; Pred. No. 7.5e+02;
tive 0; Mismatches 2;
                                                                                                     US-09-179-298-9
; Sequence 9, Application US/09179298
; Patent No. 6355421
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                477 TATCTTGGCCAGTATAT 493
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: DNA
ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: DNA ORGANISM: Artificial Sequence
17 TATATTGGCCAATATAT
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       LOCATION: (9) -
OTHER INFORMATION: spacer
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LOCATION: (9)
OTHER INFORMATION: spacer
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: misc_feature
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MOLECULE TYPE: DNA
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US-08-912-951-160
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                                                                                                                                                                                                                                                                                                                                                                                                                              RESULT 776
US-09-888-341-8
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                                                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: Cech, Thomas R.
APPLICANT: Lingner, Joachim
APPLICANT: Lingner, Joachim
APPLICANT: Lingner, Joachim
APPLICANT: Makamura, Toru
APPLICANT: Makamura, Toru
APPLICANT: Morin, Gregg B.
APPLICANT: Morin, Gregg B.
APPLICANT: Andrews, William H.
TITLE OF INVENTION: HUMAN TELOMERASE CATALYTIC SUBUNIT: DIAGNOSTIC AND
TITLE OF INVENTION: THERAPEUTIC METHODS
NUMBER OF SEQUENCES: 335
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET: Two Embarcadero Center, 8th Floor
CITY: San Francisco
STATE: California
COUNTRY: United States of America
                                                                                  ö
                                                                                  Gaps
   Score 13.8; DB 1; Length 18; Pred. No. 7.5e+02; 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/912,951
FILING DATE: US/08/912,951
FLING DATE: US/08/912,951
FLING DATE: US/08/912,951
FLING DATE: US/08/912,951
FLING DATE: US/08/912,957
CLASSIFICATION NUMBER: US 08/854,050
FILING DATE: US-MAY-1997
CLASSIFICATION: 435
FRICK APPLICATION DATA:

APPLICATION NUMBER: US 08/851,843
FILING DATE: US-MAY-1997
CLASSIFICATION: 435
FRICK APPLICATION DATA:

APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
CLASSIFICATION: 435
FRICK APPLICATION DATA:

APPLICATION NUMBER: US 08/44,419
FILING DATE: US-APR-1997
CLASSIFICATION: 435
FRICK APPLICATION DATA:

APPLICATION NUMBER: US 08/44,419
FILING DATE: US-CT-1996
CLASSIFICATION: 435
FRICK APPLICATION DATA:

APPLICATION NUMBER: US 08/44,419
FILING DATE: US-CT-1996
CLASSIFICATION: 435
FRICK APPLICATION: 435
FRICK APPLICATION UMBER: US 08/44,419
FILING DATE: US-CT-1996
CLASSIFICATION NUMBER: US-CT-1996
FLING DATE: US-CT-1996
CLASSIFICATION NUMBER: US-CT-1996
FLING DATE: US-CT-1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                      Sequence 160, Application US/08912951
Patent No. 6475789
GENERAL INFORMATION:
                                                                                                                                           477 TATCTTGGCCAGTATAT 493
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INFORMATION FOR SEQ ID NO: 160:
SEQUENCE CHARACTERISTICS:
LENGTH: 18 base pairs
TYPE: nucleic acid
STRANDEDNESS: single
   0.6%;
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Query Match 0.69
Best Local Similarity 88.29
Matches 15, Conservative
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; Sequence 8, Application US/0988341
; Patent No. 6528267
; GENERAL INFORMATION:
; APPLICANT: Coull, James M.
; APPLICANT: Coull, James M.
; APPLICANT: Hyldig-Nielsen, Jens J.
; TITLE OF INVENTION: Methods, Kits and Compositions Pertaining To PNA
; TITLE OF INVENTION: Methods, Kits and Compositions Pertaining To PNA
; TITLE OF INVENTION: Molecular Beacons
; FILE REFERENCE: BP9702US-CP1-DV1
; CURRENT APPLICATION NUMBER: US/09/888,341
; CURRENT FILING DATE: 1997-10-27
; PRIOR FILING DATE: 1997-10-27
; PRIOR FILING DATE: 1997-10-27
; PRIOR FILING DATE: 1991-10-27
; NUMBER OF SEQ ID NOS: 13
; SOFTWARE: PatentIN Ver. 2.1
; SEQ ID NO 8
                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LOCATION: (18)
OTHER INFORMATION: 3' Dabcyl
OTHER INFORMATION: Description of Artificial Sequence:SYNTHETIC PROBE
OTHER INFORMATION: OR TARGET
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Gaps
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APPLICANT: Coull, James M.
APPLICANT: Gildea, Brian D.
APPLICANT: Gildea, Brian D.
APPLICANT: Hyldig-Vielsen, Jens J.
TITLE OF INVENTION: Methods, Kits and Compositions Pertaining To PNA
TITLE OF INVENTION: Molecular Beacons
FILE REPERENCE: B9702US-CTI-DV1
CURRENT APPLICATION NUMBER: US/09/888,341
CURRENT FILING DATE: 2001-06-22
PRIOR PILING DATE: 1997-10-27
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Score 13.8; DB 1; Length 18;
Pred. No. 7.5e+02;
0; Mismatches 2; Indels
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88.2%; Pred. No. 7.5e+02;
tive 0; Mismatches 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: DNA
ORGANISM: Artificial Sequence
FEATURE:
NAME/KEY: misc_feature
LOCATION: (1)
OTHER INFORMATION: 5' Fluorescein
NAME/KEY: misc_feature
LOCATION: (9). 7(10)
OTHER INFORMATION: spacer
NAME/KEY: misc_feature
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Patent No. 6528267
                                                                                                        2279 TCACCGTGTTAGCCAGG 2295
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     0.6%;
88.2%;
                          Best Local Similarity 88.2
Matches 15, Conservative
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Matches 15; Conservative
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US-09-888-341-9
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Parent No. 6528267

GENERAL INFORMATION:
APPLICANT: Coull, James M.
APPLICANT: Gildea, Brian D.
APPLICANT: Gildea, Brian D.
APPLICANT: Hyldig-Niellen, Jens J.
APPLICANT: Hyldig-Niellen, Jens J.
APPLICANT: Hyldig-Niellen, Jens J.
TITLE OF INVENTION: Motcular Beacons
FILE REFERENCE: BP9702US-CPI-DVI
CURRENT APPLICATION NUMBER: 08/09/888,341

CURRENT FILING DATE: 1091-06-22

PRIOR FILING DATE: 10997-10-27

PRIOR FILING DATE: 1998-10-27

PRIOR FILING DATE: 1998-10-27

NUMBER: OF SEQ ID NOS: 13

SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: spacer OTHER INFORMATION: Spacer OTHER INFORMATION: Description of Artificial Sequence:SYNTHETIC PROBE OTHER INFORMATION: OR TARGET US-09-888-341-9
                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION: 3' Dabcyl
OTHER INFORMATION: Description of Artificial Sequence:SYNTHETIC PROBE
OTHER INFORMATION: OR TARGET
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match 0.6%; Score 13.8; DB 1; Length 18; Best Local Similarity 88.2%; Pred. No. 7.5e+02; Matches 15; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                         LOCATION: (1) —
OTHER INFORMATION: 5' Fluorescein NAME/KEY: misc feature
LOCATION: (9)...(10)
OTHER INFORMATION: spacer
LOCATION: (18)
LOCATION: (18)
PRIOR APPLICATION NUMBER: 09/179,298 PRIOR FILING DATE: 1998-10-27 NUMBER OF SEQ ID NOS: 13 SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        477 TATCTTGGCCAGTATAT 493
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         477 TATCTTGGCCAGTATAT 493
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                                                                                                                                                                   ORGANISM: Artificial Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: misc_feature LOCATION: (9)
                                                                                                                                                                                                                NAME/KEY: misc feature
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LENGTH: 18
                                                                                               SEQ ID NO 8
LENGTH: 18
                                                                                                                                                 TYPE: DNA
                                                                                                                                                                                                 FEATURE:
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Sequence 7339, Application US/09422978
; Sequence 7339, Application US/09422978
; Batent No. 6537751
; GENERAL INFORMATION:
; APPLICANT: Chumarkov, Inya
; TITLE OF INVENTION: Biallelic markers for use in constructing a high density...;
; TITLE OF INVENTION: Biallelic markers for use in constructing a high density...;
; TITLE OF INVENTION: Biallelic markers for use in constructing a high density...;
; TITLE OF INVENTION: Biallelic markers for use in constructing a high density...;
; TITLE OF INVENTION: Biallelic markers for use in constructing a high density...;
; TALE REPRENCE: 680872.020
; EARLIER FILING DATE: 1999-10-20
; EARLIER FILING DATE: 1999-04-21
; EARLIER APPLICATION NUMBER: US 60/109,732
; EARLIER APPLICATION NUMBER: US 60/082,614
; EARLIER FILING DATE: 1998-04-21
; NUMBER OF SEQ ID NOS: 11796
; TUMMER OF SEQ ID NOS: 11796
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Sequence 9, Application US/09888341

Patent No. 6528267

GENERAL INFORMATION:

APPLICANT: Coull, James M.

APPLICANT: Gildea, Brian D.

APPLICANT: Hyldig-Nielsen, Jens J.

TITLE OF INVENTION: Methods, Kits and Compositions Pertaining To PNA

TITLE OF INVENTION: Molecular Beacons

FILE REFERENCE: BP9702M3-CP1-DV1

CURRENT PELING DATE: 2001-06-22

PRIOR FILING DATE: 1997-10-27

PRIOR PLING DATE: 1997-10-27

PRIOR FILING DATE: 1998-10-27

NUMBER OF SEQ ID NOS: 13

SOFTWARE: PatentIn Ver. 2.1

SEQ ID NO 9

LENGTH: 18
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; OTHER INFORMATION: upstream amplification primer 99-4266 for SEQ 3464,
US-09-422-978-7398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LOCATION: (9) OTHER INFORMATION: spacer
OTHER INFORMATION: Description of Artificial Sequence:SYNTHETIC PROBE
OTHER INFORMATION: OR TARGET
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0.6%; Score 13.8; DB 1;
Best Local Similarity 88.2%; Pred. No. 7.5e+02;
Matches 15; Conservative 0; Mismatches 2;
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ORGANISM: Artificial Sequence
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Matches 15, Conservative
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ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NAME/KEY: misc_feature
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US-09-422-978-7398
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846 CGACAAAGAAAACGCCA 862

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US-09-888-341-9/c

RESULT 779

CCACAAAGAAAACTCCA 17

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Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 633
CORRESPONDENCE ADDRESS:
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COMPUTER READABLE FOOM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMpatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Townsend and Crew LLP
Center, Eighth Floor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                OTHER INFORMATION: /note= "TCP1.17 primer" SEQUENCE DESCRIPTION: SEQ ID NO: 393:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 015389-002620US
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APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIOR APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-0CT-1996
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/854,050
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/91,312
FILING DATE: 14-AQUG-1997
APPLICATION NUMBER: US 08/91,312
FILING DATE: 14-AQUG-1997
APPLICATION NUMBER: US 08/91,312
FILING DATE: 14-AQUG-1997
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AQUG-1997
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FILING DATE: 29-Sep-1997
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER: US 08/915,503
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REGISTRATION NUMBER: 42,271
                                                                                                                                                                                                           Sequence 393, Application US/09402181B Patent No. 6610839 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TELECOMMUNICATION INFORMATION: TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: Townsend and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: Two Embarcadero
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Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ATTORNEY/AGENT INFORMATION:
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INFORMATION FOR SEQ ID NO: 393:
SEQUENCE CHARACTERISTICS:
TYPE: nucleic acid
STRANDEDNESS: single
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STATE: California
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LOCATION: 1..18
OTHER INFORMATION:
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                                                                                                                                                                                                           GENERAL INFORMATION:

APPLICANT: Cohen, Daniel

APPLICANT: Cohen, Daniel

APPLICANT: Chumakov, Ilya

APPLICANT: Chumakov, Ilya

TITLE OP INVENTION: Biallelic markers for use in constructing a high density...

FILE REPERENCE: GENSET.020CP1

CURRENT APPLICATION NUMBER: US/09/422,978

CURRENT APPLICATION NUMBER: US 09/298,850

EARLIER FILING DATE: 1999-04-21

EARLIER PILING DATE: 1999-10-20

EARLIER PILING DATE: 1998-11-23

EARLIER PILING DATE: 1998-11-23

EARLIER PILING DATE: 1998-04-21

EARLIER FILING DATE: 1998-04-21

SERLIER PILING DATE: 1998-04-21
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sequence 217, Application US/09371772B
sequence 217, Application US/09371772B
sequence 217, Application US/09371772B
sequence 217, Application.
GENERAL INFORMATION:
APPLICANT: Blocky Pam Pharmaceuticals, Inc.
APPLICANT: Brocky Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Stinchcomb, Dan
APPLICANT: Brocky Jam
APPLIC
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; OTHER INFORMATION: upstream amplification primer 99-2697 for SEQ 3834,
US-09-422-978-7768
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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; Sequence 7768, Application US/09422978
; Patent No. 6537751
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Best Local Similarity 88.27
Then 15; Conservative
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ORGANISM: Homo sapiens
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ORGANISM: Homo Sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: primer_bind
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US-09-402-181B-393

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TYPE: nucleic acid
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LENGTH: 18
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                   Length 18;
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Harley, Calvin B.
Andrews, William H.
TITLE OF INVENTION: Human Telomerase Catalytic Subunit
NUMBER OF SEQUENCES: 727
CORRESPONDENCE ADDRESS:
ADDRESSEE: Townsend and Townsend and Crew LLP
STREET Two Embarcadero Center, Eighth Floor
CITY: San Francisco
                                                        2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ZIF: 9411-3834

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 09-MAY-1997
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/911,312
APPLICATION NUMBER: US 08/912,951
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: US 08/915,503
FILING DATE: 14-AUG-1997
APPLICATION NUMBER: WO PCT/US97/17618
FILING DATE: 01-OCT-1997
APPLICATION NUMBER: WO PCT/US97/17885
FILING DATE: 01-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    REFERENCE/DOCKET NUMBER: 015389-002610US
               Score 13.8; DB 1;
Pred. No. 7.5e+02;
0; Mismatches 2;
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APPLICATION NUMBER: US/08/974,549A
FILING DATE: 19.00V-1997
APPLICATION NUMBER: US 08/724,643
FILING DATE: 01-0CT-1996
APPLICATION NUMBER: US 08/844,419
FILING DATE: 18-APR-1997
APPLICATION NUMBER: US 08/846,017
FILING DATE: 25-APR-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
APPLICATION NUMBER: US 08/851,843
FILING DATE: 06-MAY-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/09/721,456
FILING DATE: 22-No. 6617110-2000
CLASSIFICATION: <UNKNOWN>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NAME: Apple, Randolph Ted
REGISTRATION NUMBER: 36,429
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 576-0200
                                                                                                                                                                                                                                        Sequence 393, Application US/09721456
Patent No. 6617110
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (415) 576-0300
                                                                                                                                                                                                                                                                                                       APPLICANT: Cech, Thomas R.
Lingner, Joachim
Nakamura, Toru
Chapman, Karen B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                 2279 TCACCGTGTTAGCCAGG 2295
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               INFORMATION FOR SEQ ID NO: 393: SEQUENCE CHARACTERISTICS:
                                                                                                                      2 TCACCGTGTTGGGCAGG 18
            0.6%;
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                                                        15; Conservative
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                                  Similarity
                                                                                                                                                                                                  RESULT 784
US-09-721-456-393
            Query Match
Best Local {
                                                        Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ; OTHER INFORMATION: Description of Artificial Sequence: Primer US-09-465-491-4
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ROSSAU, RUDI
MAENTENS, GEERT
TITLE OF INVENTION: METHOD FOR TYPING AND DETECTING HBV
NUMBER OF SEQUENCES: 313
CORRESPONDENCE ADDRESS:
ADDRESSEE: NIXON & VANDERHYE P.C.
STREET: 1100 NORTH GLEBE ROAD
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                                                                                                                                                                                 Length 18;
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                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 4. Application US/09465491
Patent No. 6664046
GENERAL INFORMATION:
APPLICANT: Chang, Sheng-Yung P.
APPLICANT: Cantini, Christopher D.
TITLE OF INVENTION: Quantitation of hTERT mRNA Expression; FILE REFERENCE: RPA1002
CURRENT APPLICATION NUMBER: US/09/465,491
CURRENT FILING DATE: 1999-12-16
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver: 2.1
                             0.6%; Score 13.8; DB 1;
88.2%; Pred. No. 7.5e+02;
tive 0; Mismatches 2;
                                                                                                                                                                               0.6%; Score 13.8; DB 1;
88.2%; Pred. No. 7.5e+02;
tive 0; Mismatches 2;
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Patent No. 6709812
GENERAL INFORMATION:
APPLICANT: STUYVER, LIEVEN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U.S.A.
ZIP: 22201-4714
COMPUTER READABLE FORM
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                 2279 TCACCGTGTTAGCCAGG 2295
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                                                                                                                                                                                                                                                                                                          rcaccererresecase 18
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ORGANISM: Artificial Sequence
TOPOLOGY: linear MOLECULE TYPE: DNA
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Best Local Similarity 88.2
Matches 15; Conservative
                                                                                                                                                                               Query Match
Best Local Similarity 88.2
Matches 15; Conservative
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Search completed: January 25, 2005, 15:32:19 Job time : 28 Becs